



SEQUENCE LISTING

<110> Nixon, Andrew
Hogan, Shannon

<120> PAPP-A LIGANDS

<130> 10280-059001

<140> US 10/783,311

<141> 2004-02-19

<150> US 60/448,515

<151> 2003-02-19

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<170> FastSEQ for Windows Version 4.0

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<213> Homo sapiens

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Ser	Leu	Glu	Pro	Pro	Leu	Cys	Gly	Gln	Thr	Leu	Cys	Asp	Asn	Thr	Glu	
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Val	Ile	Ala	Ser	Tyr	Asn	Gln	Leu	Ser	Phe	Arg	Gln	Pro	Lys	Val		
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Cys	Tyr	Leu	Asp	Leu	Val	Tyr	Gln	Gly	Trp	Gln	Pro	Ser	Arg	Lys	Pro	
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Ala	Pro	Val	Ala	Leu	Ala	Pro	Gln	Val	Leu	Gly	His	Thr	Thr	Asp	Ser	
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Gln Tyr Ala Ser Asn Ala Ser Ser Pro Met Pro Cys Ser Pro Ser Gly				
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His Trp Ser Pro Arg Glu Ala Glu Gly His Pro Asp Val Glu Gln Pro				
		740		745
Cys Lys Ser Ser Val Arg Thr Trp Ser Pro Asn Ser Ala Val Asn Pro				
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His Thr Val Pro Pro Ala Cys Pro Glu Pro Gln Gly Cys Tyr Leu Glu				
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Leu Glu Phe Leu Tyr Pro Leu Val Pro Glu Ser Leu Thr Ile Trp Val				
785		790		795
Thr Phe Val Ser Thr Asp Trp Asp Ser Ser Gly Ala Val Asn Asp Ile				
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Lys Leu Leu Ala Val Ser Gly Lys Asn Ile Ser Leu Gly Pro Gln Asn				
		820		825
Val Phe Cys Asp Val Pro Leu Thr Ile Arg Leu Trp Asp Val Gly Glu				
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Glu Val Tyr Gly Ile Gln Ile Tyr Thr Leu Asp Glu His Leu Glu Ile				
		850		855
Asp Ala Ala Met Leu Thr Ser Thr Ala Asp Thr Pro Leu Cys Leu Gln				
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Cys Lys Pro Leu Lys Tyr Lys Val Val Arg Asp Pro Pro Leu Gln Met				
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Asp Val Ala Ser Ile Leu His Leu Asn Arg Lys Phe Val Asp Met Asp				
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Leu Asn Leu Gly Ser Val Tyr Gln Tyr Trp Val Ile Thr Ile Ser Gly				
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Thr Glu Glu Ser Glu Pro Ser Pro Ala Val Thr Tyr Ile His Gly Arg				
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Gly Tyr Cys Gly Asp Gly Ile Ile Gln Lys Asp Gln Gly Glu Gln Cys				
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Asp Asp Met Asn Lys Ile Asn Gly Asp Gly Cys Ser Leu Phe Cys Arg				
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Gln Glu Val Ser Phe Asn Cys Ile Asp Glu Pro Ser Arg Cys Tyr Phe				
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His Asp Gly Asp Gly Val Cys Glu Glu Phe Glu Gln Lys Thr Ser Ile				
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Lys Asp Cys Gly Val Tyr Thr Pro Gln Gly Phe Leu Asp Gln Trp Ala				
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Ser Asn Ala Ser Val Ser His Gln Asp Gln Gln Cys Pro Gly Trp Val				
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Ile Ile Gly Gln Pro Ala Ala Ser Gln Val Cys Arg Thr Lys Val Ile				
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Asp Leu Ser Glu Gly Ile Ser Gln His Ala Trp Tyr Pro Cys Thr Ile				
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Ser Tyr Pro Tyr Ser Gln Leu Ala Gln Thr Thr Phe Trp Leu Arg Ala				
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Tyr Phe Ser Gln Pro Met Val Ala Ala Ala Val Ile Val His Leu Val				
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Thr Asp Gly Thr Tyr Tyr Gly Asp Gln Lys Gln Glu Thr Ile Ser Val				
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Gln Leu Leu Asp Thr Lys Asp Gln Ser His Asp Leu Gly Leu His Val				
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Leu Ser Cys Arg Asn Asn Pro Leu Ile Ile Pro Val Val His Asp Leu				
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				1150

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 Pro Leu Val Ala Ile Ser Gly Val Ala Leu Arg Ser Phe Asp Asn Phe
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 Asp Pro Val Thr Leu Ser Ser Cys Gln Arg Gly Glu Thr Tyr Ser Pro
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 Glu Leu Ala Val Glu Asn Ala Ser Leu Asn Cys Ser Ser Ser Asp Arg
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 Tyr His Gly Ala Gln Cys Thr Val Ser Cys Arg Thr Gly Tyr Val Leu
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 Lys Cys Glu Asp Ser Asp Ala Ser Gln Gly Leu Gly Ser Asn Val Ile
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 His Cys Arg Lys Asp Gly Thr Trp Asn Gly Ser Phe His Val Cys Gln
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 Lys Leu Gln Cys Pro Asp Gly Tyr Ala Ile Gly Ser Glu Cys Ala Thr
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 Val Val Cys Thr Ala Gly Leu Lys Trp Tyr Pro His Pro Ala Leu Ile
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 His Cys Val Lys Gly Cys Glu Pro Phe Met Gly Asp Asn Tyr Cys Asp
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 Thr Ser Thr Val Lys Thr Lys Lys Val Thr Pro Phe Pro Met Ser Cys
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 Asp Leu Gln Gly Asp Cys Ala Cys Arg Asp Pro Gln Ala Gln Glu His

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Glu	Leu	Pro	Arg	Asp	Ala	Phe	Thr	Leu	Gln	Val	Trp	Leu	Arg	Ala	Glu				
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Ser	Tyr	Ile	Ser	Arg	Asp	Arg	Gly	Trp	Val	Val	Gly	Ile	His	Thr	Ile				
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Pro	Gly	Gln	Trp	Val	Tyr	Leu	Ala	Ala	Thr	Tyr	Asp	Gly	Gln	Phe	Met				
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Gly	Gly	Ser	Ala	Leu	Asn	His	Asn	Tyr	Arg	Gly	Tyr	Ile	Glu	His	Phe				
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Ser	Leu	Trp	Lys	Val	Ala	Arg	Thr	Gln	Arg	Glu	Ile	Leu	Ser	Asp	Met				
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Glu	Thr	His	Gly	Ala	His	Thr	Ala	Leu	Pro	Gln	Leu	Leu	Gln	Glu					
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Val	Ile	Ala	Ser	Tyr	Asn	Gln	Leu	Ser	Ser	Phe	Arg	Gln	Pro	Lys	Val				
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Val	Arg	Tyr	Arg	Val	Val	Asn	Leu	Tyr	Glu	Asp	Asp	His	Lys	Asn	Pro				
		275					280					285							
Thr	Val	Thr	Arg																

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Arg Phe Asn Phe Asp Gly	Gly Glu Cys Cys Asp	Pro Glu Ile Thr Asn
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Val Thr Gln Thr Cys Phe	Asp Pro Asp Ser Pro	His Arg Ala Tyr Leu
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Asp Val Asn Glu Leu Lys	Asn Ile Leu Lys Leu	Asp Gly Ser Thr His
420	425	430
Leu Asn Ile Phe Phe Ala	Lys Ser Ser Glu Glu Glu	Leu Ala Gly Val
435	440	445
Ala Thr Trp Pro Trp Asp	Lys Glu Ala Leu Met	His Leu Gly Gly Ile
450	455	460
Val Leu Asn Pro Ser Phe	Tyr Gly Met Pro Gly	His Thr His Thr Met
465	470	475
Ile His Glu Ile Gly His	Ser Leu Gly Leu Tyr	His Val Phe Arg Gly
485	490	495
Ile Ser Glu Ile Gln Ser	Cys Ser Asp Pro Cys	Met Glu Thr Glu Pro
500	505	510
Ser Phe Glu Thr Gly Asp	Leu Cys Asn Asp Thr	Asn Pro Ala Pro Lys
515	520	525
His Lys Ser Cys Gly Asp	Pro Gly Pro Gly Asn	Asp Thr Cys Gly Phe
530	535	540
His Ser Phe Phe Asn Thr	Pro Tyr Asn Asn Phe	Met Ser Tyr Ala Asp
545	550	555
Asp Asp Cys Thr Asp Ser	Phe Thr Pro Asn Gln	Val Ala Arg Met His
565	570	575
Cys Tyr Leu Asp Leu Val	Tyr Gln Gly Trp Gln	Pro Ser Arg Lys Pro
580	585	590
Ala Pro Val Ala Leu Ala	Pro Gln Val Leu Gly	His Thr Thr Asp Ser
595	600	605
Val Thr Leu Glu Trp Phe	Pro Pro Ile Asp Gly	His Phe Phe Glu Arg
610	615	620
Glu Leu Gly Ser Ala Cys	His Leu Cys Leu Glu	Gly Arg Ile Leu Val
625	630	635
Gln Tyr Ala Ser Asn Ala	Ser Ser Pro Met Pro	Cys Ser Pro Ser Gly
645	650	655
His Trp Ser Pro Arg Glu	Ala Glu Gly His Pro	Asp Val Glu Gln Pro
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Cys Lys Ser Ser Val Arg	Thr Trp Ser Pro Asn	Ser Ala Val Asn Pro
675	680	685
His Thr Val Pro Pro Ala	Cys Pro Glu Pro Gln	Gly Cys Tyr Leu Glu
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Leu Glu Phe Leu Tyr Pro	Leu Val Pro Glu Ser	Leu Thr Ile Trp Val
705	710	715
Thr Phe Val Ser Thr Asp	Trp Asp Ser Ser Gly	Ala Val Asn Asp Ile
725	730	735
Lys Leu Leu Ala Val Ser	Gly Lys Asn Ile Ser	Leu Gly Pro Gln Asn
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Val Phe Cys Asp Val Pro	Leu Thr Ile Arg Leu	Trp Asp Val Gly Glu
755	760	765
Glu Val Tyr Gly Ile Gln	Ile Tyr Thr Leu Asp	Glu His Leu Glu Ile
770	775	780
Asp Ala Ala Met Leu Thr	Ser Thr Ala Asp Thr	Pro Leu Cys Leu Gln
785	790	795
Cys Lys Pro Leu Lys Tyr	Lys Val Val Arg Asp	Pro Pro Leu Gln Met
805	810	815
Asp Val Ala Ser Ile Leu	His Leu Asn Arg Lys	Phe Val Asp Met Asp
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 Thr Glu Glu Ser Glu Pro Ser Pro Ala Val Thr Tyr Ile His Gly Arg
 850 855 860
 Gly Tyr Cys Gly Asp Gly Ile Ile Gln Lys Asp Gln Gly Glu Gln Cys
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 Asp Asp Met Asn Lys Ile Asn Gly Asp Gly Cys Ser Leu Phe Cys Arg
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 Asp Leu Ser Glu Gly Ile Ser Gln His Ala Trp Tyr Pro Cys Thr Ile
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 Ser Tyr Pro Tyr Ser Gln Leu Ala Gln Thr Thr Phe Trp Leu Arg Ala
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 Tyr Phe Ser Gln Pro Met Val Ala Ala Ala Val Ile Val His Leu Val
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 Ala Glu Gln Ser Cys Val His Phe Ala Cys Glu Lys Thr Asp Cys Pro
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 Glu Leu Ala Val Glu Asn Ala Ser Leu Asn Cys Ser Ser Ser Asp Arg
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 Tyr His Gly Ala Gln Cys Thr Val Ser Cys Arg Thr Gly Tyr Val Leu
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 Gln Ile Arg Arg Asp Asp Glu Leu Ile Lys Ser Gln Thr Gly Pro Ser
 1170 1175 1180
 Val Thr Val Thr Cys Thr Glu Gly Lys Trp Asn Lys Gln Val Ala Cys
 1185 1190 1195 1200
 Glu Pro Val Asp Cys Ser Ile Pro Asp His His Gln Val Tyr Ala Ala
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 Ser Phe Ser Cys Pro Glu Gly Thr Thr Phe Gly Ser Gln Cys Ser Phe
 1220 1225 1230
 Gln Cys Arg His Pro Ala Gln Leu Lys Gly Asn Asn Ser Leu Leu Thr
 1235 1240 1245
 Cys Met Glu Asp Gly Leu Trp Ser Phe Pro Glu Ala Leu Cys Glu Leu
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<223> Light Chain nucleic acid

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ctctcctgca	gggccagtca	gagtgttagt	aggaacttag	cctggtacca	gcagaaacct	180
ggccaggctc	ccaggctcct	catctatggt	gcatccacca	gggccactgg	tatcccagcc	240
aggttcagt	gcagtgggtc	tgggacagag	ttcactctca	ccatcagcag	cctgcagtct	300
gaagattttg	cagtttatca	ctgtcagcag	tataatagca	ggcctctcac	tttcggcgga	360
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ctctcctgca	gggccagtca	gagtgttcgc	agctacttag	cctggtacca	gcagaaacca	180
ggccaggctc	ccaggctcct	catctatgat	gcatccacca	gggccactgg	tatcccagcc	240
agattcagt	gcagtgggtc	tgggacagag	ttcactctca	ccatcagcag	cctgcagtct	300
gaagattttg	cagtttatta	ctgtcagcag	tataataact	ggcctccgac	gttcggccaa	360
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ctctcctgca	gggccagtca	ggatgttaac	agatacttag	cctggtacca	gcagaaacct	180
ggccagcctc	ccaggctcct	catctatggt	gcctctacca	gggccactgg	tatcccagcc	240
aggatcagt	gcagtgggtc	tgggacagag	ttcactctca	ccatcagcag	cctgcagtct	300
gaagattttg	cagtttatta	ctgtcagcag	tatcataact	ggcccctcac	tttcggcgga	360
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gtgaaaaaat	tattattcgc	aattccttta	gttggttcctt	tctattctca	cagtgcacaa	60
gacatccaga	tgacccagtc	tccagccacc	ctgtctttgt	ctccagggga	aagagccacc	120

ctctcctgca	gggccagtca	gagtgttagc	agctacttag	cctggtacca	acagaaacct	180
ggccaggctc	ccaggctcct	catctatggt	gcatccagca	gggccactgg	catcccagac	240
aggttcagt	gcagtgggtc	tgggacagac	ttcactctca	ccatcggcag	actggagcct	300
gaagattttg	cagtgtatta	ctgtcagcag	tatagtagtt	caccggtcac	cttcggccaa	360
gggacacgac	tggagattaa	acgaactgtg	gctgcaccat	ctgtc		405

<210> 8

<211> 402

<212> DNA

<213> Artificial Sequence

<220>

<223> Light Chain nucleic acid sequence

<400> 8

gtgaaaaaat	tattattcgc	aattccttta	gttggttcctt	tctattctca	cagtgcacaa	60
gacatccaga	tgaccagtc	tccagccacc	ctgtctttgt	ctccagggga	aagagccacc	120
ctctcctgca	gggccagtca	gagtgttagc	aggtacttag	cctggtacca	acagaaacct	180
ggccaggctc	ccaggctcct	catctatggt	gcatccacca	gggccactgg	tatcccagcc	240
aggttcagt	gcagtgggtc	tgggacagag	ttcactctca	ccatcagcag	cctgcagtct	300
gaagattttg	cagtttatta	ctgtcagcag	tataataact	ggccttcttt	cggcggaggg	360
accaaggtgg	agatcaaacg	aactgtggct	gcaccatctg	tc		402

<210> 9

<211> 408

<212> DNA

<213> Artificial Sequence

<220>

<223> Light Chain nucleic acid sequence

<400> 9

gtgaaaaaat	tattattcgc	aattccttta	gttggttcctt	tctattctca	cagtgcacaa	60
gacatccaga	tgaccagtc	tccaggcacc	ctgtctttgt	ctccagggga	aagagccacc	120
ctctcctgca	gggccagtca	gagtattagc	agcagctact	tagcctggta	ccagcagaaa	180
cctggccagg	ctcccaggct	cctcatctat	gctgcagcca	gcagggccac	tggcatccca	240
gacaggttca	gtggcattgg	gtctgggaca	gacttcactc	tcaccatcag	cagcctagag	300
cctgaagatt	ttgcagttta	ttactgtcag	cagcgtagca	actggcctct	cactttcggc	360
ggagggacca	aggtggagat	caaacgaact	gtggctgcac	catctgtc		408

<210> 10

<211> 408

<212> DNA

<213> Artificial Sequence

<220>

<223> Light Chain nucleic acid sequence

<400> 10

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gacatccaga	tgaccagtc	tccaggcacc	ctgtctttgt	ctccagggga	aagagccacc	120
ctctcctgca	gggccagtca	gagtgttagc	agcagctact	tagcctggta	ccagcagaaa	180
cctggccagg	ctcccaggct	cctcatctat	ggtgcatcca	gcagggccac	tggcatccca	240
gacaggttca	gtggcagtgg	gtctgggaca	gacttcactc	tcaccatcag	cagactggag	300
cctgaagatt	ttgcagtgtg	ttactgtcag	cagtatggta	gctcaccgtg	gacgttcggc	360
caagggacca	aggtggaaaat	caaacgaact	gtggctgcac	catctgtc		408

<210> 11
 <211> 405
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Light Chain nucleic acid sequence

<400> 11
 gtgaaaaaat tattattcgc aattccttta gttgttcctt tctattctca cagtgcacaa 60
 gacatccaga tgaccagtc tccatcctca ctgtctgcat ctgtaggaga cagagtcacc 120
 atcacttgtc gggcgagtc ggacattagc aattatttag cctggtttca gcagaaacca 180
 gggagagccc ctaagtcctt gatctatggt gcatccagtt tgcaaactgg ggtcccatca 240
 aagttcagcg gcagtggatc tgggacagag ttactctca ccatcagcgg cctgcagcct 300
 gaagatgttg caacttatta ctgccatcag tataatcatt accctcccac tttcggcgga 360
 gggaccaagg tggagatcaa acgaactgtg gctgcacat ctgtc 405

<210> 12
 <211> 405
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Light Chain nucleic acid sequence

<400> 12
 gtgaaaaaat tattattcgc aattccttta gttgttcctt tctattctca cagtgcacaa 60
 gacatccaga tgaccagtc tccatcctca ctgtctgcat ctgtaggaga cagagtcacc 120
 atcacttgtc gggcgagtc ggacattagc aattatttag cctggtttca gcagaaacca 180
 ggggaagccc ctaagtcctt gatctatgct ggtccagtt tgcaagtggt ggtctcatca 240
 aacttcagcg gcagtggatc tgggacagat ttactctca ccatcagcag cctgcagcct 300
 gaagattttg caacttatta ctgccagcag tatcataggt acccgaggac ttttggtcag 360
 gggaccaagc tggagatcaa acgaactgtg gctgcacat ctgtc 405

<210> 13
 <211> 560
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Light Chain nucleic acid sequence

<400> 13
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 gacatccaga tgaccagtc tccatcctca ttctctgcat ctacaggaga cagagtcacc 120
 atcacttgtc gggcgagtc gggattagc agttatttag cctggatatca gcaaaaacca 180
 gggaaagccc ctaagtcctt gatctatgct gcatccactt tgcaaagtgg ggtcccatca 240
 aagttcagcg gcagtggatc tgggacagat ttactctca ccatcagcag cctgcagcct 300
 gaagattttg caacttatta ctgccaacag tataatagtt accccctcac cttcggccaa 360
 gggacacgac tggagattaa acgaactgtg gctgcacat ctgtcttcat cttcccgcca 420
 tctgatgagc agttgaaatc tggaactgcc tctgttgtgt gcctgctgaa taacttctat 480
 cccagagagg ccaaagtaca gtggaagggt gataacgccc tccaatcggg taactccag 540
 gagagtgtca cagagcagga 560

<210> 14
 <211> 405
 <212> DNA

<213> Artificial Sequence

<220>

<223> Light Chain nucleic acid sequence

<400> 14

gtgaaaaaat	tattattcgc	aattccttta	gttggtcctt	tctattctca	cagtgcacaa	60
gacatccaga	tgacccagtc	tccatcctcc	ctgtatgcat	ctgtaggaga	cagagtcacc	120
atcacttgcc	gggcaagtca	gggcattaga	aatgagttag	gttggtatca	gcagaaacca	180
gggaaagccc	ctcagcgcct	gatctatgat	gcatccactt	tgcaagtggt	gggtcccatca	240
agattcagcg	gcggtggatc	taggacagaa	ttcactctca	ccatcagcag	cctggaacct	300
catgattttg	gaacttatta	ctgccaacaa	tatgccagtt	atccgctcac	tttcggcgga	360
gggaccaagg	tggagatcaa	acgaactgtg	gctgcaccat	ctgtc		405

<210> 15

<211> 405

<212> DNA

<213> Artificial Sequence

<220>

<223> Light Chain nucleic acid sequence

<400> 15

gtgaaaaaat	tattattcgc	aattccttta	gttggtcctt	tctattctca	cagtgcacaa	60
gacatccaga	tgacccagtc	tccatcctcc	ctgtctgcat	ctgtaggaga	cagagtcacc	120
atcacttgcc	gggcaagtca	gagcattagc	agctatttaa	attggtatca	gcagaaacca	180
gggaaagccc	ctaagctcct	gatctatgct	gcatccagtt	tgcaaagtgg	gggtcccatca	240
aggttcagtg	gcagtggatc	tgggacagat	ttcactctca	ccatcagcag	tctgcaacct	300
gaagattttg	caacttacta	ctgtcaacag	agttacagta	ccaggtggac	gttcggccaa	360
gggaccaagg	tggaaatcaa	acgaactgtg	gctgcaccat	ctgtc		405

<210> 16

<211> 405

<212> DNA

<213> Artificial Sequence

<220>

<223> Light Chain nucleic acid sequence

<400> 16

gtgaaaaaat	tattattcgc	aattccttta	gttggtcctt	tctattctca	cagtgcacaa	60
gacatccaga	tgacccagtc	tccatcctcc	ctgtctgcat	ctgtaggaga	cagagtcacc	120
atcacttgcc	gggcaagtca	gagcattagc	agctatttaa	attggtatca	gcagaaacca	180
gggaaagccc	ctaagctcct	gatctatgct	gcatccagtt	tgcaaagtgg	gggtcccatca	240
aggttcagtg	gcagtggatc	tgggacagat	ttcactctca	ccatcagcag	tctgcaacct	300
gaagattttg	caacttacta	ctgtcaacag	agttacagta	ccaggtggac	gttcggccaa	360
gggaccaagg	tggaaatcaa	acgaactgtg	gctgcaccat	ctgtc		405

<210> 17

<211> 560

<212> DNA

<213> Artificial Sequence

<220>

<223> Light Chain nucleic acid sequence

<400> 17

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gtgaaaaaat tattattcgc aattccttta gttgttcctt tctattctca cagtgcacaa      60
gacatccaga tgacccagtc tccatcctcc ctgtctgcat ctgtaggaga cagagtcacc      120
atcacttgcc gggcaagtca gggcattaga aatgatttag gctgggttca gcagaaacca      180
gggaaagccc ctaggcgctt gatctggggt gcatccactt taaaaagtgg ggtcccacatca    240
aggttcagcg gcagtggatc tggcacagat ttcactctca ccatcagcag cctgcagcct      300
gaagattttg caacttatta ctgtctacaa gattacaatt acccgtagac ttttggccag      360
gggaccaagc tggagatcaa acgaactgtg gctgcaccat ctgtcttcat cttcccgcga      420
tctgatgagc agttgaaatc tggaaactgc tctgttgtgt gcctgctgaa taacttctat      480
cccagagagg ccaaagtaca gtggaagggt gataacgcc tccaatcggg taactcccag      540
gagagtgtca cagagcagga

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<210> 18

<211> 405

<212> DNA

<213> Artificial Sequence

<220>

<223> Light Chain nucleic acid sequence

<400> 18

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gtgaaaaaat tattattcgc aattccttta gttgttcctt tctattctca cagtgcacaa      60
gacatccaga tgacccagtc tccatcctcc ctgtctgcat ctgtaggaga cagagtcacc      120
atcacttgcc gggcaagtca gggcattaga cattatttag gctgggtatca gcagaaacca      180
gggaaagccc ctaagcgctt gatctatgct gcatccagtt tgcaatttgg ggtcccagca      240
aggttcagcg gcagtggatc tgggacggaa ttcactctca caatcagcag cctgcagcct      300
gaagattttg caacttatta ctgtctacaa cacaatagtt tccctccggc gttcggccaa      360
gggaccaagg tggaaatcaa acgaactgtg gctgcaccat ctgtc

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<210> 19

<211> 405

<212> DNA

<213> Artificial Sequence

<220>

<223> Light Chain nucleic acid sequence

<400> 19

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gtgaaaaaat tattattcgc aattccttta gttgttcctt tctattctca cagtgcacaa      60
gacatccaga tgacccagtc tccatcctcc ctgtctgcat ctgtaggaga cagagtcacc      120
atcacttgcc gggcaagtca gggcattaga cattatttag gctgggtatca gcagaaacca      180
gggaaagccc ctaagcgctt gatctatgct gcatccagtt tgcaatttgg ggtcccagca      240
aggttcagcg gcagtggatc tgggacggaa ttcactctca caatcagcag cctgcagcct      300
gaagattttg caacttatta ctgtctacaa cacaatagtt tccctccggc gttcggccaa      360
gggaccaagg tggaaatcaa acgaactgtg gctgcaccat ctgtc

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<210> 20

<211> 405

<212> DNA

<213> Artificial Sequence

<220>

<223> Light Chain nucleic acid sequence

<400> 20

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gtgaaaaaat tattattcgc aattccttta gttgttcctt tctattctca cagtgcacaa      60
gacatccaga tgacccagtc tccatcctcc ctgtctgcat ctgtaggaga cagagtcacc      120
atcacttgcc gggcaagtca gggcattaga cattatttag gctgggtatca gcagaaacca      180

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gggaaagccc	ctaagcgcct	gatctatgct	gcatccagtt	tgcaatttgg	ggccccagca	240
aggttcagcg	gcagtggatc	tgggacggaa	ttcactctca	caatcagcag	cctgcagcct	300
gaagattttg	caacttatta	ctgtctacaa	cacaatagtt	tccctccggc	gttcggccaa	360
gggaccaagg	tggaaatcaa	acgaactgtg	gctgcaccat	ctgtc		405

<210> 21

<211> 405

<212> DNA

<213> Artificial Sequence

<220>

<223> Light Chain nucleic acid sequence

<400> 21

gtgaaaaaat	tattattcgc	aattccttta	gttggttcctt	tctattctca	cagtgcacaa	60
gacatccaga	tgaccagtc	tccatcctcc	ctgtctgcat	ctgtaggaga	cagagtcacc	120
atctcttgcc	gcgcaagtca	gaacattagg	aactctgtaa	attggtatca	gcagaaacca	180
gggaaagccc	ctaagctcct	gatctatgct	acatacgatt	tgcaagtggt	cgccccatca	240
tacttcagtg	gcagtggatc	tgggacagat	ttcactctca	ccatcaccag	tctgcaacct	300
gaagattttg	caacttacta	ctgtcaacag	agttacagtt	tccctcgaac	gttcggccaa	360
gggaccaagg	tggaaatcag	acgaactgtg	gctgcaccat	ctgtc		405

<210> 22

<211> 405

<212> DNA

<213> Artificial Sequence

<220>

<223> Light Chain nucleic acid sequence

<400> 22

gtgaaaaaat	tattattcgc	aattccttta	gttggttcctt	tctattctca	cagtgcacaa	60
gacatccaga	tgaccagtc	tccatcttcc	gtgtctgcat	ctgtaggaga	cagaatcgcc	120
atcacttgtc	gggcgagtc	gggtattagc	acctgggttag	cctgggtatca	gcagagacca	180
gggagagccc	ctaagctcct	gatctatgct	gcatccactt	tgcaaaagcg	agtcccatca	240
aggttcagcg	gcagtggatc	tgggacagat	ttcactctca	ccatcagcag	cctgcagcct	300
gaagattttg	caacttactt	ttgtcaacag	gctgacagtt	tccccctgac	ttttggccag	360
gggaccaaac	tgagatcaa	acgaactgtg	gctgcaccat	ctgtc		405

<210> 23

<211> 405

<212> DNA

<213> Artificial Sequence

<220>

<223> Light Chain nucleic acid sequence

<400> 23

gtgaaaaaat	tattattcgc	aattccttta	gttggttcctt	tctattctca	cagtgcacaa	60
gacatccaga	tgaccagtc	tccatcttcc	gtgtctgcat	ctgtaggaga	cagagtcacc	120
atcacttgtc	gggcgagtc	gggtattagc	agatgggttag	cctgggtatca	gcagaaacca	180
gggaaagccc	ctaagctcct	gatctatggt	gcatccactt	tgcaaaaagg	ggccccatca	240
aggttcaccg	gcagtggatc	tgggacagat	ttcactctca	ccatcaccag	cctgcagcct	300
gaagattttg	caacttacta	ttgtcaacag	ggtaacagtt	tcccattcac	tttcggccct	360
gggaccaaag	tggatatcaa	acgaactgtg	gctgcaccat	ctgtc		405

<210> 24

<211> 405
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Light Chain nucleic acid sequence

<400> 24
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 gacatccaga tgaccagtc tccatcttcc gtgtctgcat ctgtaggaga cagagtcacc 120
 atcacttgtc gggcgagtcg ggggtattagc agatgggttag cctgggtatca gcagaaacca 180
 gggaaagccc ctaagctcct gatctatggt gcatccactt tgcaaaaagg ggtcccatca 240
 aggttcaccg gcagtggatc tgggacagat ttcactctca ccatcaccag cctgcagcct 300
 gaagattttg caacttacta ttgtcaacag ggtaacagtt tcccattcac tttcggccct 360
 gggaccaaaag tggatatcaa acgaactgtg gctgcaccat ctgtc 405

<210> 25
 <211> 405
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Light Chain nucleic acid sequence

<400> 25
 gtgaaaaaat tattattcgc aattccttta gttgttcctt tctattctca cagtgcacaa 60
 gacatccaga tgaccagtc tccgtcttcc gtgtctgcat ctgtaggaga cagagtcacc 120
 atcacttgtc gggcgagtca ggggtattagc agctgggttag cctgggtatca gcagaagcca 180
 gggaaagccc ctaagttgct gatctatggt gcatccagtt tggaaagtgg ggtcccatca 240
 agattcagcg gcagtggatc tgggacagat tacactctca ccatcaccag cctacagcct 300
 gaagattttg caacttactt ttgtcaacag gttaattctt tccctcgtac ttttggccag 360
 gggaccaagc tgaatatcaa acgaactgtg gctgcaccat ctgtc 405

<210> 26
 <211> 539
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Light Chain nucleic acid sequence

<400> 26
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 agcgaattga ctgagataag ggccagggtc ctggagtccc cagccgcttc tctggatcca 120
 aagatgctgc agccaatgca ggggttttac tcatctccgg cctccagccc gaggatgatg 180
 ctgactatct ttgtatgata tgggttaagca atgtacatgc gacattcggc ggagggacca 240
 agctgaccgt cctgggtcag cccaaggctg cccctcgggt cactctgttc ccgccctcct 300
 ctgaggagct tcaagccaac aaggccacac tgggtgtgtc cataagtgc ttctaccggg 360
 gagccgtgac agtggcctgg aaggcagata gcagccccgt caaggcggga gtggagacca 420
 ccacaccctc caaacaagc aacaacaagt acgcgccag cagctatcta agcctgacgc 480
 ctgagcagtg gaagtccac agaagctaca gctgccaggt cacgcatgaa gggagcacc 539

<210> 27
 <211> 411
 <212> DNA
 <213> Artificial Sequence

<220>

<223> Light Chain nucleic acid sequence

<400> 27

gtgaaaaaat	tattattcgc	aattccttta	gttggttcctt	tctattctca	cagtgcacag	60
agcgaattga	ctcagccacc	ctcagcgtct	gcgacccccg	ggcagagggg	caccatctct	120
tgttctggaa	gcagctccaa	catcggacgt	aatttggtat	actggtacca	gcagctccca	180
ggaacggccc	ccaaactcct	catctatagt	aataatcagc	ggccctcagg	ggccctgac	240
cgattctctg	gctccaagtc	tggcacctca	gcctccctgg	ccatcagtgg	gctccgggtcc	300
gaggaggagg	ctgattatta	ctgtgcagca	tgggatgaca	gcctgagtgg	ttgggtgttc	360
ggcggagggg	ccaggctgac	cgtcctaggt	cagcccaagg	ctgccccctc	g	411

<210> 28

<211> 411

<212> DNA

<213> Artificial Sequence

<220>

<223> Light Chain nucleic acid sequence

<400> 28

gtgaaaaaat	tattattcgc	aattccttta	gttggttcctt	tctattctca	cagtgcacag	60
agcgctttga	ctcagccacc	ctcagcgtct	gggacccccg	ggcagagggg	caccatctct	120
tgttctggaa	gcagctccaa	catcgggaagt	aattttgtat	actggtacca	ccatctccca	180
ggaacggccc	ccaaactcct	catctatagg	aataatcagc	ggccctcagg	ggccctgac	240
cgattctctg	gctccaagtc	tggcacctca	gcctccctgg	ccatcagtgg	gctccgggtcc	300
gaggatgagg	ctgattatta	ctgtgcagca	tgggatgaca	gcctgagtgg	gggtggtattc	360
ggcggagggg	ccaagctgac	cgtcctaggt	cagcccaagg	ctgccccctc	g	411

<210> 29

<211> 414

<212> DNA

<213> Artificial Sequence

<220>

<223> Light Chain nucleic acid sequence

<400> 29

gtgaaaaaat	tattattcgc	aattccttta	gttggttcctt	tctattctca	cagtgcacag	60
agcgctttga	ctcagcctgc	ctccgtgtct	gggtctcctg	gacagtcgat	caccatctcc	120
tgcactggaa	ccagcagtga	cgttgggtat	tatgactatg	tctcctggta	ccagcaccac	180
ccaggcaaag	ccccaaact	catcatttat	gatgtcactt	ctcggccctc	aggggtctct	240
tctcatttct	ctggctccaa	gtctggcaac	acggcctccc	tgaccatctc	tgggtccag	300
gctgatgacg	aggctgatta	ttactgcagc	tcatatacaa	gcggcagcac	ccgttatgtc	360
ttcggacctg	ggaccaaggt	caccgtccta	ggtcagccca	aggccaacc	cact	414

<210> 30

<211> 414

<212> DNA

<213> Artificial Sequence

<220>

<223> Light Chain nucleic acid sequence

<400> 30

gtgaaaaaat	tattattcgc	aattccttta	gttggttcctt	tctattctca	cagtgcacag	60
agcgctttga	ctcagcctgc	ctccgtgtct	gggtctcctg	gacagtcgat	caccatctcc	120

tgcactggaa	ccagcagtga	cgttgggtat	tatgactatg	tctcctggta	ccagcaccac	180
ccaggcaaag	cccccaaact	catcatttat	gatgtcactt	ctcggccctc	aggggtctct	240
tctcatttct	ctggctccaa	gtctggcaac	acggcctccc	tgaccatctc	tgggtccag	300
gctgatgacg	aggctgatta	ttactgcagc	tcatatacaa	gcggcagcac	ccgttatgtc	360
ttcggacctg	ggaccaaggt	caccgtccta	ggtcagccca	aggccaaccc	caact	414

<210> 31

<211> 414

<212> DNA

<213> Artificial Sequence

<220>

<223> Light Chain nucleic acid sequence

<400> 31

gtgaaaaaat	tattattcgc	aattccttta	gttggtcctt	tctattctca	cagtgcacag	60
agcgtcttga	ctcagactgc	ctccgtgtct	gggtctcctg	gacagtcgat	caccatctcc	120
tgcactggaa	ccagcagtga	cattgggtgat	tatgagtatg	tctcctggta	ccaacaacac	180
ccaggcaaag	cccccaaagt	cattcctttat	gaggtcagta	atcggccctc	aggggtccct	240
gatcgcttct	ctggctccaa	gtctggcaac	acggcctcac	tgaccatctc	tggactccag	300
gctgaggacg	aggctgatta	ttactgtggt	tcatatagaa	agagcagcac	tccttatgtc	360
ttcggaaactg	ggaccaaggt	cagcgtccta	ggtcagccca	aggccaaccc	caact	414

<210> 32

<211> 414

<212> DNA

<213> Artificial Sequence

<220>

<223> Light Chain nucleic acid sequence

<400> 32

gtgaaaaaat	tattattcgc	aattccttta	gttggtcctt	tctattctca	cagtgcacag	60
agcgtcttga	ctcagactgc	ctccgtgtct	gggtctcctg	gacagtcgat	caccatctcc	120
tgcactggaa	ccagcagtga	cattgggtgat	tatgagtatg	tctcctggta	ccaacaacac	180
ccaggcaaag	cccccaaagt	cattcctttat	gaggtcagta	atcggccctc	aggggtccct	240
gatcgcttct	ctggctccaa	gtctggcaac	acggcctcac	tgaccatctc	tggactccag	300
gctgaggacg	aggctgatta	ttactgtggt	tcatatagaa	agagcagcac	tccttatgtc	360
ttcggaaactg	ggaccaaggt	cagcgtccta	ggtcagccca	aggccaaccc	caact	414

<210> 33

<211> 560

<212> DNA

<213> Artificial Sequence

<220>

<223> Light Chain nucleic acid sequence

<400> 33

gtgaaaaaat	tattattcgc	aattccttta	gttggtcctt	tctattctca	cagtgcacag	60
agcgtcttga	ctcagccacc	ctcagcgtct	gggacccccg	ggcagagggg	caccatctct	120
tgttctggaa	gcagctccaa	catcgaaagt	aatactgtaa	cctggtacca	gcaactccca	180
ggaacggccc	ccaaactcct	catctatagt	gatgatcagc	ggccctcagg	ggtccctgac	240
cgattctctg	gatccaagtc	tggcacctca	gcctccctgg	ccatcagtgg	gctccagtct	300
gaggatgagg	ctgactatta	ctgtgcaaca	tgggataaca	ccctgagagg	tgtgggtttc	360
ggcggaggga	ccaagctgac	cgtcctgagt	cagcccaagg	ctgccccctc	ggtcactctg	420
ttccccccct	cctctgagga	gcttcaagcc	aacaaggcca	caactggtgtg	tctcataagt	480

gacttctacc cgggagccgt gacagtggcc tggaaggcag atagcagccc cgtcaaggcg 540
ggagtggaga ccaccacacc 560

<210> 34

<211> 429

<212> DNA

<213> Artificial Sequence

<220>

<223> Light Chain nucleic acid sequence

<400> 34

gtgaaaaaat	tattattcgc	aattccttta	gttggttcctt	tctattctca	cagtgcacag	60
agcgtcttga	ctcagccacc	ttatgcatca	gcctccctgg	gagcctcgg	cacactcacc	120
tgcaccctga	gcagcggcta	cagtaattat	aaagtggact	ggatcagca	aagaccaggg	180
aagggccccc	agtttgtgat	gcgagtgggc	agtggcggga	ttgtgggatc	aaagggggat	240
ggcatccctg	atcgcttttc	agtcctgggc	tcaggcctgt	atcggtatct	gaccatcaag	300
aacatccagg	aagaggatga	gagtgactac	tattgtgggg	cagaccatgg	cagggggggc	360
accttcgtgt	gggtgttcgg	cggagggacc	aaactgaccg	tcctaggtca	gccaaggct	420
gccccctcg						429

<210> 35

<211> 411

<212> DNA

<213> Artificial Sequence

<220>

<223> Light Chain nucleic acid sequence

<400> 35

gtgaaaaaat	tattattcgc	aattccttta	gttggttcctt	tctattctca	cagtgcacag	60
agcgtcttga	ctcagcctgc	ctccgtgtct	gggtctcctg	gacagtcgat	caccatctcc	120
tgcactggaa	ccagcagtga	cgttgggtgt	tataactatg	tctcctggta	ccaacgacac	180
ccaggcaaag	cccccaaact	cattatttat	gatgtcacta	atcgccctc	aggggttct	240
cgtcattct	ctggctccaa	gtctggcaac	acggcctccc	tgaccatctc	tggtctccag	300
gccgacgacg	aggctgatta	ttattgcgtt	tcatttacaa	acagcaatac	tttcgtcttc	360
ggaagtggga	ccagggtcac	cgtcctcgg	cagcccaagg	ccaacccac	t	411

<210> 36

<211> 417

<212> DNA

<213> Artificial Sequence

<220>

<223> Light Chain nucleic acid sequence

<400> 36

gtgaaaaaat	tattattcgc	aattccttta	gttggttcctt	tctattctca	cagtgcacag	60
gacatcgtca	tgactcaaac	ccctcctagt	ttaccggtta	acccgggtga	acctgcctcc	120
atctcctgca	agtctagtca	gagcctcctg	cagagtaatg	gatacaacta	cttggattgg	180
tacctgcaga	agccagggca	gtctccacag	ctcctgatct	atctgggttc	taatcggggc	240
tccgggggtcc	ctgacaggtt	cagtggcagt	ggatcaggca	cagattttac	actgaagatc	300
agcaggggtg	aggctgagga	tggtggcatt	tattactgca	tgcaagctct	acacactcct	360
cccttcggcc	aagggtacac	actggagatt	aaacgaactg	tggctgcacc	atctgtc	417

<210> 37

<211> 405

<212> DNA

<213> Artificial Sequence

<220>

<223> Light Chain nucleic acid sequence

<400> 37

gtgaaaaaat	tattattcgc	aattccttta	gttgttcctt	tctattctca	cagtgcacag	60
tacgaattga	ctcagccacc	ctcagtgtcc	gtgtccccgg	gacagacagc	caccattatc	120
tgctctggag	ataaattggg	ggataaatat	gttgcctggg	atcagcagaa	gccaggccag	180
tcccctgtgc	tggtcgtcta	tgaagataac	aagcggccct	cagggatccc	tgagcgaatt	240
tctggctcca	actctgggaa	cacagccact	ctgaccatca	gtgggaccca	ggctatggat	300
gacgctgact	attactgtca	ggcgtgggac	agaagcactg	accattatgt	cttcggaact	360
gggaccaagg	tcaccgtcct	aggtcagccc	aaggccaacc	ccact		405

<210> 38

<211> 414

<212> DNA

<213> Artificial Sequence

<220>

<223> Light Chain nucleic acid sequence

<400> 38

gtgaaaaaat	tattattcgc	aattccttta	gttgttcctt	tctattctca	cagtgcacag	60
tacgaattga	ctcagcctgc	ctccgtgtct	gggtctcctg	gacagtcat	caccatctcc	120
tgcaactggaa	ccagcagcga	cggtgggtgg	tataactatg	tctcctggta	ccaacagcac	180
ccaggcaaag	ccccaaact	catgatttat	gaggtcagta	atcgccctc	aggggtttct	240
aatcgcttct	ctggctccaa	gtctgacaat	acggcctccc	tgaccatctc	tggactccag	300
gctgaggacg	aggctgatta	ttactgtggg	tcatatagaa	agagcagcac	tccttatgtc	360
ttcggaaactg	ggaccaaggt	cagcgtccta	ggtcagccca	aggccaaccc	cact	414

<210> 39

<211> 413

<212> DNA

<213> Artificial Sequence

<220>

<223> Light Chain nucleic acid sequence

<400> 39

gtgaaaaaat	tatttattcg	caatttcctt	tagttgttcc	tttctattct	cacagtgcac	60
agagcgcttt	gactcagcca	tcctcagcgt	ctgggacccc	cgggcagagg	gtcagtatct	120
cttgttctgg	aagcagctac	aacatcggag	tttatgatgt	atactggtac	cagcagctcc	180
caggaacggc	ccccaaactc	ctcatctata	ccaataatca	gcggccctca	ggggtccctg	240
accgattctc	tggtccaag	tctggcacct	cagcctccct	ggccatcagt	gggctccagt	300
ctgaggatga	ggctgattat	tactgtgcag	catgggatga	cagtctgagt	ggttgggtgt	360
tcggcggagg	gaccaaggtg	accgtcctag	gtcagcccaa	ggctgcccc	tcg	413

<210> 40

<211> 387

<212> DNA

<213> Artificial Sequence

<220>

<223> Heavy Chain nucleic acid sequence

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<400> 40
gaagttcaat tgtagagtc tggtagcggt cttgttcagc ctggtggttc tttacgtctt      60
tcttgcgctg cttccggatt cactttctct ctttaccaga tgctttgggt tcgccaagct      120
cctggtaaag gtttggagtg ggtttctggt atcgtttctt ctggtggcct tactggttat      180
gctgactccg ttaaaggctg cttcactatc tctagagaca actctaagaa tactctctac      240
ttgcagatga acagcttaag ggctgaggac actgcagtct actattgtgc gagacataat      300
agggtattg gcacctttga ctactggggc cagggaaccc tggtcaccgt ctcaagcgcc      360
tccaccaagg gccatcggt cttcccg

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<210> 41
<211> 369
<212> DNA
<213> Artificial Sequence

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<220>
<223> Heavy Chain nucleic acid sequence

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<400> 41
gaagttcaat tgtagagtc tggtagcggt cttgttcagc ctggtggttc tttacgtctt      60
tcttgcgctg cttccggatt cactttctct tggtagctta tgaattgggt tcgccaagct      120
cctggtaaag gtttggagtg ggtttcttct atctatcctt ctggtggcta tactatgtat      180
gctgactctg ttaaaggctg cttcactatc tctagagaca actctaagaa gactctctac      240
ttgcagatga acagcttaag ggctgaggac actgcagtct actattgtgc gagtgacttt      300
ggtagctggg gccaggggaa cctggtcacc gtctcaagcg cctccaccaa gggcccatcg      360
gtcttcccg

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<210> 42
<211> 369
<212> DNA
<213> Artificial Sequence

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<220>
<223> Heavy Chain nucleic acid sequence

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<400> 42
gaagttcaat tgtagagtc tggtagcggt cttgttcagc ctggtggttc tttacgtctt      60
tcttgcgctg cttccggatt cactttctct tggtagccta tgaattgggt tcgccaagct      120
cctggtaaag gtttggagtg ggtttcttct atcgttcctt ctggtggcta tactcgttat      180
gctgactccg ttaaaggctg cttcactatc tctagagaca actctaagaa tactctctac      240
ttgcagatga acagcttaag ggctgaggac actgcagtct actattgtgc gagtgacttt      300
ggtagctggg gccaggggaa cctggtcacc gtctcaagcg cctccaccaa gggcccatcg      360
gtcttcccg

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<210> 43
<211> 369
<212> DNA
<213> Artificial Sequence

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<220>
<223> Heavy Chain nucleic acid sequence

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<400> 43
gaagttcaat tgtagagtc tggtagcggt cttgttcagc ctggtggttc tttacgtctt      60
tcttgcgctg cttccggatt cactttctct cgttactcta tgaattgggt tcgccaagct      120
cctggtaaag gtttggagtg ggtttcttat atctctcctt ctggtggcat gactaagtat      180
gctgactccg ttaaaggctg cttcactatc tctagagaca actctaagaa tactctctac      240
ttgcagatga acagcttaag ggctgaggac actgcagtct actattgtgc gaataccctt      300

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ggctactggg gccaggggaac cctgggtcacc gtctcaagcg cctccaccaa gggcccatcg 360
gtcttccccg 369

<210> 44

<211> 369

<212> DNA

<213> Artificial Sequence

<220>

<223> Heavy Chain nucleic acid sequence

<400> 44

gaagttcaat	tgtagagtc	tggtggcgg	cttggtcagc	ctggtggttc	tttacgtctt	60
tcttgcgctg	cttccggatt	cactttctct	tcttaccgta	tgaattgggt	tcgccaagct	120
cctggtaaaag	gtttggagtg	ggtttctggt	atcggttcctt	ctggtggcaa	gactttttat	180
gctgactccg	ttaaagggtcg	cttcactatc	tctagagaca	actctaagaa	tactctctac	240
ttgcagatga	acagcttaag	ggctgaggac	actgcagtct	actattgtgc	gagtgacttt	300
ggtagctggg	gccaggggaac	cctgggtcacc	gtctcaagcg	cctccaccaa	gggcccatcg	360
gtcttccccg						369

<210> 45

<211> 369

<212> DNA

<213> Artificial Sequence

<220>

<223> Heavy Chain nucleic acid sequence

<400> 45

gaagttcaat	tgtagagtc	tggtggcgg	cttggtcagc	ctggtggttc	tttacgtctt	60
tcttgcgctg	cttccggatt	cactttctct	aattactcta	tggattgggt	tcgccaagct	120
cctggtaaaag	gtttggagtg	ggtttcttgg	atctctcctt	ctggtggcct	tactacttat	180
gctgactccg	ttaaagggtcg	cttcactatc	tctagagaca	actctaagaa	tactctctac	240
ttgcagatga	acagcttaag	ggctgaggac	actgcagtct	actattgtgc	gagtgacttt	300
ggtagctggg	gccaggggaac	cctgggtcacc	gtctcaagcg	cctccaccaa	gggcccatcg	360
gtcttccccg						369

<210> 46

<211> 393

<212> DNA

<213> Artificial Sequence

<220>

<223> Heavy Chain nucleic acid sequence

<400> 46

gaagttcaat	tgtagagtc	tggtggcgg	cttggtcagc	ctggtggttc	tttacgtctt	60
tcttgcgctg	cttccggatt	cactttctct	cgttaccata	tggagtgggt	tcgccaagct	120
catggtaaaag	gtttggagtg	ggtttcttat	atctctcctt	ctggtggcaa	gactctttat	180
gctgactccg	ttaaagggtcg	cttcactatc	tctagagaca	actctaagaa	tactctctac	240
ttgcagatga	acagcttaag	ggctgaggac	actgcagtct	actattgtgc	gagacatttg	300
ggatatgggt	cggggagtta	ctttgactac	tggggccagg	gaaccctggg	caccgtctca	360
agcgcctcca	ccaagggccc	atcgggtcttc	ccg			393

<210> 47

<211> 390

<212> DNA

<213> Artificial Sequence

<220>

<223> Heavy Chain nucleic acid sequence

<400> 47

gaagttcaat	tgtagagtc	tggtggcgg	cttggtcagc	ctggtggttc	tttacgtctt	60
tcttgcgctg	cttccggatt	cactttctct	ttttacccta	tgcttggtg	tcgccaagct	120
cctggtaaag	gtttgagtg	ggtttcttat	atctctcctt	ctggtggcga	tactacttat	180
gctgactccg	ttaaaggctg	cttcactatc	tctagagaca	actctaagaa	tactttctac	240
ttgcagatga	acagcttaag	ggctgaggac	actgcagtct	actattgtgc	gagagggggg	300
tcctatagca	gcagttggta	cggctactgg	ggccagggaa	ccctggtcac	cgtctcaagc	360
gcctccacca	agggcccatc	ggtcttcccg				390

<210> 48

<211> 387

<212> DNA

<213> Artificial Sequence

<220>

<223> Heavy Chain nucleic acid sequence

<400> 48

gaagttcaat	tgtagagtc	tggtggcgg	cttggtcagc	ctggtggttc	tttacgtctt	60
tcttgcgctg	cttccggatt	cactttctct	aagtacccta	tgcttggtg	tcgccaagct	120
cctggtaaag	gtttgagtg	ggtttcttgg	atctctcctt	ctggtggcaa	gactgtttat	180
gctgactccg	ttaaaggctg	cttcactatc	tctagagaca	actctaagaa	tactctctac	240
ttgcagatga	acagcttaag	ggctgaggac	actgcagtct	actattgtgc	gaaagattgc	300
agaggggggt	gcagtggtgg	aagttggggc	caggggaacc	tggtcacctg	ctcaagcgcc	360
tccaccaagg	gcccacgggt	cttcccg				387

<210> 49

<211> 393

<212> DNA

<213> Artificial Sequence

<220>

<223> Heavy Chain nucleic acid sequence

<400> 49

gaagttcaat	tgtagagtc	tggtggcgg	cttggtcagc	ctggtggttc	tttacgtctt	60
tcttgcgctg	cttccggatt	cactttctct	gcttacaata	tgcttggtg	tcgccaagct	120
cctggtaaag	gtttgagtg	ggtttcttat	atctcttctt	ctggtactgg	ttatgctgac	180
tccgttaaag	gtcgcttcac	tatctctaga	gacaactcta	agaatactct	ctacttgacg	240
atgaacagct	taagggctga	ggacactgca	gtctactatt	gtgcgagaga	actgggtagt	300
gggagctact	acccgggata	cttccagcac	tggggccagg	gcaccctggt	caccgtctca	360
agcgctcca	ccaagggccc	atcggtcttc	ccg			393

<210> 50

<211> 369

<212> DNA

<213> Artificial Sequence

<220>

<223> Heavy Chain nucleic acid sequence

<400> 50

gaagttcaat	tgtagagtc	tggtggcgg	cttggtcagc	ctggtgggtc	tttacgtctt	60
tcttgcgctg	cttccggatt	cactttctct	tggtacacta	tggtttgggt	tcgccaagct	120
cctggtaaag	gtttggagtg	ggtttcttct	atctattctt	ctggtgggtt	tacttggtat	180
gctgactccg	ttaaaggctg	cttcactatc	tctagagaca	actctaagaa	tactctctac	240
ttgcagatga	acagcttaag	ggctgaggac	actgcagtct	actattgtgc	gagtgacttt	300
ggtagctggg	gccagggaac	cctggtcacc	gtctcaagcg	cctccaccaa	gggcccatcg	360
gtcttccc						369

<210> 51

<211> 420

<212> DNA

<213> Artificial Sequence

<220>

<223> Heavy Chain nucleic acid sequence

<400> 51

gaagttcaat	tgtagagtc	tggtggcgg	cttggtcagc	ctggtgggtc	tttacgtctt	60
tcttgcgctg	cttccggatt	cactttctct	gattacaaga	tgcttgggt	tcgccaagct	120
cctggtaaag	gtttggagtg	ggtttcttct	atctgggtctt	ctggtggcac	tactgagtat	180
gctgactccg	ttaaaggctg	cttcactatc	tctagagaca	actctaagaa	tactctctac	240
ttgcagatga	acagcttaag	ggctgaggac	actgcagtct	actattgtgc	gagagaggaa	300
attggacgat	attttgactg	gtttttagg	aactactact	actacggtat	ggacgtctgg	360
ggccaaggga	ccacggtcac	cgtctcaagc	gcctccacca	agggcccatc	ggtcttccc	420

<210> 52

<211> 411

<212> DNA

<213> Artificial Sequence

<220>

<223> Heavy Chain nucleic acid sequence

<400> 52

gaagttcaat	tgtagagtc	tggtggcgg	cttggtcagc	ctggtgggtc	tttacgtctt	60
tcttgcgctg	cttccggatt	cactttctct	acttacttta	tgcttgggt	tcgccaagct	120
cctggtaaag	gtttggagtg	ggtttcttat	atcggttcctt	ctggtggcaa	tactctttat	180
gctgactccg	ttaaaggctg	cttcactatc	tctagagaca	actctaagaa	tactctctac	240
ttgcagatga	acagcttaag	ggctgaggac	actgcagtct	actattgtgc	aagagaagag	300
tgggacgtat	tactatggtt	cggggagtta	agtgtctgctt	ttgatatctg	gggccaagg	360
acaatggtca	ccgtctcaag	cgctccacc	aagggcccat	cggtcttccc	g	411

<210> 53

<211> 369

<212> DNA

<213> Unknown

<220>

<223> Light Chain nucleic acid sequence

<400> 53

ttctattctc	acagtgcaca	gagcgaattg	actcagccac	cctcagcgtc	tgggaccccc	60
gggcagaggg	tcaccatctc	ttgttctgga	agcagctcca	acatcggaag	taatactgta	120
aactggtagc	agcagctccc	aggaacggcc	cccaaactcc	tcatctatag	taataattac	180
cggccctcag	gggtccctga	ccgattctct	ggctccaagt	ctggcacctc	agcctccctg	240
gccatcagtg	ggctccagtc	tgacgatgag	gctgaatatc	tctgtgcagc	atgggatgac	300

agtctgaatg gtccggtggt cggtggaggg accaaggtga ccgtcctagg tcagcccaag 360
gctgcccc 369

<210> 54

<211> 396

<212> DNA

<213> Artificial Sequence

<220>

<223> Heavy Chain nucleic acid sequence

<400> 54

gaagttcaat	tgtagagtc	tggtggcgg	cttggtcagc	ctggtgggtc	tttacgtctt	60
tcttgcgctg	cttcggatt	cactttctct	ttttacggta	tgccttgggt	tcgccaagct	120
cctggtaaag	gtttgagtg	ggtttctggt	atctatcctt	ctggtggcgt	tactcgttat	180
gctgactccg	ttaaaggctg	cttcactatc	tctagagaca	actctaagaa	tactctctac	240
ttgcagatga	acagcttaag	ggctgaggac	actgcagtct	actattgtgc	gaagacgtat	300
agcagcagct	ggtacgggtg	gtactttgac	tactggggcc	aggggaaccct	ggtcaccgtc	360
tcaagcgcct	ccaccaaggg	cccatcggtc	ttcccc			396

<210> 55

<211> 396

<212> DNA

<213> Artificial Sequence

<220>

<223> Heavy Chain nucleic acid sequence

<400> 55

gaagttcaat	tgtagagtc	tggtggcgg	cttggtcagc	ctggtgggtc	tttacgtctt	60
tcttgcgctg	cttcggatt	cactttctct	ccttacgata	tgtggtgggt	tcgccaagct	120
cctggtaaag	gtttgagtg	ggtttcttat	atctcttctt	ctggtggcaa	gactatgtat	180
gctgactccg	ttaaaggctg	cttcactatc	tctagagaca	actctaagaa	tactctctac	240
ttgcagatga	acagcttaag	ggctgaggac	actgcagtct	actattgtgc	gagattaggt	300
ggtaactccc	actactacta	cggtatggac	gtctggggcc	aagggaaccac	ggtcaccgtc	360
tcaagcgcct	ccaccaaggg	cccatcggtc	ttcccc			396

<210> 56

<211> 396

<212> DNA

<213> Artificial Sequence

<220>

<223> Heavy Chain nucleic acid sequence

<400> 56

gaagttcaat	tgtagagtc	tggtggcgg	cttggtcagc	ctggtgggtc	tttacgtctt	60
tcttgcgctg	cttcggatt	cactttctct	ccttacgata	tgtggtgggt	tcgccaagct	120
cctggtaaag	gtttgagtg	ggtttcttat	atctcttctt	ctggtggcaa	gactatgtat	180
gctgactccg	ttaaaggctg	cttcactatc	tctagagaca	actctaagaa	tactctctac	240
ttgcagatga	acagcttaag	ggctgaggac	actgcagtct	actattgtgc	gagattaggt	300
ggtaactccc	actactacta	cggtatggac	gtctggggcc	aagggaaccac	ggtcaccgtc	360
tcaagcgcct	ccaccaaggg	cccatcggtc	ttcccc			396

<210> 57

<211> 390

<212> DNA

<213> Unknown

<220>

<223> Heavy Chain nucleic acid sequence

<400> 57

gaagttcaat	tgtagagtc	tggtggcgg	cttggtcagc	ctggtgggtc	tttacgtctt	60
tcttgcgctg	cttccggatt	cactttctct	tcttacgtta	tgatttgggt	tcgccaagct	120
cctggtaaag	gtttgagtg	ggtttcttg	atctcttctt	ctggtggcta	tacttcttat	180
gctgactccg	ttaaaggtcg	cttcactatc	tctagagaca	actctaagaa	tactctctac	240
ttgcagatga	acagcttaag	ggctgaggac	actgcagtct	actactgtgc	gaaagggccc	300
gggacccggg	gtgactactg	gggccagga	accctgggtca	ccgtctcaag	cgctccacc	360
aagggcccat	cggtcttccc	gctagcacc				390

<210> 58

<211> 351

<212> DNA

<213> Artificial Sequence

<220>

<223> Heavy Chain nucleic acid sequence

<400> 58

gaagttcaat	tgtagagtc	tggtggcgg	cttggtcagc	ctggtgggtc	tttacgtctt	60
tcttgcgctg	cttccggatt	cactttctct	ccttacacta	tgaattgggt	tcgccaagct	120
cctggtaaag	gtttgagtg	ggtttctcg	atcggttctt	ctggtgttta	ctcattatgc	180
tgactccggt	aaaggtcgct	tactatctc	tagagacaac	tctaagaata	ctctctactt	240
gcagatgaac	agcttaaggg	ctgaggacac	tgcagtctac	tattgtgcga	gaccaccct	300
ctattggtat	ggttcgggga	gctattacta	ctttgactac	tggggccagg	g	351

<210> 59

<211> 369

<212> DNA

<213> Artificial Sequence

<220>

<223> Heavy Chain nucleic acid sequence

<400> 59

gaagttcaat	tgtagagtc	tggtggcgg	cttggtcagc	ctggtgggtc	tttacgtctt	60
tcttgcgctg	cttccggatt	cactttctct	aattacgcta	tggattgggt	tcgccaagct	120
cctggtaaag	gtttgagtg	ggtttcttat	atctctcctt	ctggtggcta	tactcgttat	180
gctgactccg	ttaaaggtcg	cttcactatc	tctagagaca	actctaagaa	tactctctac	240
ttgcagatga	acagcttaag	ggctgaggac	actgcagtct	actattgtgc	gagtgacttt	300
ggtagctggg	gccaggaac	cctggtcacc	gtctcaagcg	cctccaccaa	gggcccatcg	360
gtcttccc						369

<210> 60

<211> 396

<212> DNA

<213> Artificial Sequence

<220>

<223> Heavy Chain nucleic acid sequence

<400> 60

gaagttcaat	tgtagagtc	tggtggcgg	cttggtcagc	ctggtgggtc	tttacgtctt	60
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tcttgcgctg	cttccggatt	cactttctct	ggttactgga	tgtcttgggt	tcgccaagct	120
cctggtaaag	gtttggagtg	ggtttctgtt	atccgtcctt	ctgggtggcaa	gactgggttat	180
gctgactccg	ttaaaggctg	cttcactatc	tctagagaca	actttaagaa	tactctctac	240
ttgcagatga	acagcttaag	ggctgaggac	actgcagtct	actattgtgc	gagagtaagg	300
gcgcccggct	actactacta	cggtatggac	gtctggggcc	aagggaccac	ggtcaccgtc	360
tcaagcgct	ccaccaaggg	cccatcggtc	ttcccc			396

<210> 61

<211> 396

<212> DNA

<213> Artificial Sequence

<220>

<223> Heavy Chain nucleic acid sequence

<400> 61

gaagttcaat	tgtagagtc	tggtggcggt	cttggttcagc	ctgggtgggtc	tttacgtctt	60
tcttgcgctg	cttccggatt	cactttctct	ggttactgga	tgtcttgggt	tcgccaagct	120
cctggtaaag	gtttggagtg	ggtttctgtt	atccgtcctt	ctgggtggcaa	gactgggttat	180
gctgactccg	ttaaaggctg	cttcactatc	tctagagaca	actttaagaa	tactctctac	240
ttgcagatga	acagcttaag	ggctgaggac	actgcagtct	actattgtgc	gagagtaagg	300
gcgcccggct	actactacta	cggtatggac	gtctggggcc	aagggaccac	ggtcaccgtc	360
tcaagcgct	ccaccaaggg	cccatcggtc	ttcccc			396

<210> 62

<211> 351

<212> DNA

<213> Artificial Sequence

<220>

<223> Heavy Chain nucleic acid sequence

<400> 62

gaagttcaat	tgtagagtc	tggtggcggt	cttggttcagc	ctgggtgggtc	tttacgtctt	60
tcttgcgctg	cttccggatt	cactttctct	tcttacctta	tgacttgggt	tgccaagctc	120
ctggtaaaag	tttgagtg	gtttcttcta	tctatccttc	tggtggccat	actgggttatg	180
ctgactccgt	ttaaaggctgc	ttcactatct	ctagagacaa	ctctaagaat	actctctact	240
tgtagatgaa	cagcttaagg	gctgaggaca	ctgcagtcta	ctatttgtgcg	agagaggggg	300
gatattgtag	tagtaccagc	tgctatgttg	actactgggg	ccaggggaacc	c	351

<210> 63

<211> 414

<212> DNA

<213> Artificial Sequence

<220>

<223> Heavy Chain nucleic acid sequence

<400> 63

gaagttcaat	tgtagagtc	tggtggcggt	cttggttcagc	ctgggtgggtc	tttacgtctt	60
tcttgcgctg	cttccggatt	cactttctct	cgttacggta	tgaagtgggt	tcgccaagct	120
cctggtaaag	gtttggagtg	ggtttcttat	atctatcctt	ctgggtggcta	tactcgttat	180
gctgactccg	ttaaaggctg	cttcactatc	tctagagaca	actctaagaa	tactctctac	240
ttgcagatga	acagcttaag	ggctgaggac	actgcagtct	actattgtgc	gagagcccgc	300
gggcatagca	gcagctggta	caatcattac	tactactact	acatggacgt	ctgggggcaaa	360
gggaccacgg	tcaccgtctc	aagcgcctcc	accaagggcc	catcggtctt	cccg	414

<210> 64
 <211> 393
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Heavy Chain nucleic acid sequence

<400> 64
 gaagttcaat tgtagagtc tggtagcggt cttgttcagc ctggtgggtc tttacgtctt 60
 tcttgcgctg cttccggatt cactttctct tggtagcata tgcgttgggt tcgccaagct 120
 cctggtaaag gtttggagtg ggtttctatc tatccttctg gtggcggtac ttcttatgct 180
 gactccgtta aaggctcgtt cactatctct agagacaact ctaagaatac tctctacttg 240
 cagatgaaca gcttaagggc tgaagacact gcagtctact attgtgctgag agaaacaagt 300
 ggctgggtata gggatcgctg gttagacccc tggggccagg gaaccctggt caccgtctca 360
 agcgctcca ccaagggccc atcgggtcttc ccg 393

<210> 65
 <211> 384
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Heavy Chain nucleic acid sequence

<400> 65
 gaagttcaat tgtagagtc tggtagcggt cttgttcagc ctggtgggtc tttacgtctt 60
 tcttgcgctg cttccggatt cactttctct cagtacaaga tgaattgggt tcgccaagct 120
 cctggtaaag gtttggagtg ggtttcttat atctctcctt ctggtggcta tactgcttat 180
 gctgactccg ttaaaggctg cttcactatc tctagagaca actctaagaa tactctctac 240
 ttgcagatga acagcttaag ggctgaggac actgcagtct actattgtgc gagagatgta 300
 gtggctgggc cgtttgacta ctggggccag ggaaccctgg tcaccgtctc aagcgctccc 360
 accaagggcc catcgggtctt cccg 384

<210> 66
 <211> 393
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Heavy Chain nucleic acid sequence

<400> 66
 gaagttcaat tgtagagtc tggtagcggt cttgttcagc ctggtgggtc tttacgtctt 60
 tcttgcgctg cttccggatt cactttctct gattactata tgcgttgggt tcgccaagct 120
 cctggtaaag gtttggagtg ggtttctcgt atctatcctt ctggtggcca tacttggtat 180
 gctgactccg ttaaaggctg cttcactatc tctagagaca actctaagaa tactctctac 240
 ttgcagatga acagcttaag ggctgaggac actgcagtct actattgtgc gagacatagg 300
 gcgggtagca gtggctggta ctctgactac tggggccagg gaaccctggt caccgtctca 360
 agcgctcca ccaagggccc atcgggtcttc ccg 393

<210> 67
 <211> 393
 <212> DNA
 <213> Artificial Sequence

<220>

<223> Heavy Chain nucleic acid sequence

<400> 67

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cctggtaaag	gtttggagtg	ggtttctcgt	atctatcctt	ctggtggcca	tacttggtat	180
gctgactccg	ttaaagggtcg	cttcactatc	tctagagaca	actctaagaa	tactctctac	240
ttgcagatga	acagcttaag	ggctgaggac	actgcagtct	actattgtgc	gagacatagg	300
gcgggtagca	gtggctggta	ctctgactac	tggggccagg	gaaccctggg	caccgtctca	360
agcgctcca	ccaagggcc	atcggctctc	ccg			393

<210> 68

<211> 414

<212> DNA

<213> Artificial Sequence

<220>

<223> Heavy Chain nucleic acid sequence

<400> 68

gaagttcaat	tgtttagagtc	tggtggcggt	cttggttcagc	ctggtgggttc	tttacgtctt	60
tcttgcgctg	cttccggatt	cactttctct	tattaccata	tgtggtgggt	tcgccaagct	120
cctggtaaag	gtttggagtg	ggtttctggt	atcggtcctt	ctggtggcgg	tactcagtat	180
gctgactccg	ttaaagggtcg	cttcactatc	tctagagaca	actctaagaa	tactctctac	240
ttgcagatga	acagcttaag	ggctgaggac	actgcagtct	actattgtgc	gagagatgga	300
catagcagca	gctggtacgg	tgggggagcc	cactactacg	gtatggacgt	ctggggccaa	360
gggaccacgg	tcaccgtctc	aagcgctcc	accaagggcc	catcgggtctt	ccc	414

<210> 69

<211> 414

<212> DNA

<213> Artificial Sequence

<220>

<223> Heavy Chain nucleic acid sequence

<400> 69

gaagttcaat	tgtttagagtc	tggtggcggt	cttggttcagc	ctggtgggttc	tttacgtctt	60
tcttgcgctg	cttccggatt	cactttctct	tattaccata	tgtggtgggt	tcgccaagct	120
cctggtaaag	gtttggagtg	ggtttctggt	atcggtcctt	ctggtggcgg	tactcagtat	180
gctgactccg	ttaaagggtcg	cttcactatc	tctagagaca	actctaagaa	tactctctac	240
ttgcagatga	acagcttaag	ggctgaggac	actgcagtct	actattgtgc	gagagatgga	300
catagcagca	gctggtacgg	tgggggagcc	cactactacg	gtatggacgt	ctggggccaa	360
gggaccacgg	tcaccgtctc	aagcgctcc	accaagggcc	catcgggtctt	ccc	414

<210> 70

<211> 396

<212> DNA

<213> Artificial Sequence

<220>

<223> Heavy Chain nucleic acid sequence

<400> 70

gaagttcaat	tgtttagagtc	tggtggcggt	cttggttcagc	ctggtgggttc	tttacgtctt	60
tcttgcgctg	cttccggatt	cactttctct	ccttaccgta	tggattgggt	tcgccaagct	120
cctggtaaag	gtttggagtg	ggtttcttat	atctatcctt	ctggtggcctt	tactccttat	180

gctgactccg	ttaaagggtcg	cttcactatc	tctagagaca	actctaagaa	tactttctac	240
ttgcagatga	acagcttaag	ggctgaggac	actgcagtct	actattgtgc	gaaagggtca	300
acgggatacc	gctactacta	cggtatggac	gtctggggcc	aaggggaccac	ggtcaccgtc	360
tcaagcgcct	ccaccaaggg	cccatcggtc	ttcccc			396

<210> 71

<211> 408

<212> DNA

<213> Artificial Sequence

<220>

<223> Heavy Chain nucleic acid sequence

<400> 71

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tcttgcgctg	cttccggatt	cactttctct	tacaagatga	tgtgggttcg	ccaagctcct	120
ggtaaagggt	tggagtgggt	ttcttatatc	tcttcttctg	gtggcattac	tacttatgct	180
gactccgtta	aaggtcgctt	cactatctct	agagacaact	ctaagaatac	tctctacttg	240
cagatgaaca	gcttaagggc	tgaggacact	gcagtctact	attgtgagag	agacccgact	300
tacgattttt	ggagtgggta	ttactactac	tactacatgg	acgtctgggg	caaagggacc	360
acggtcaccg	tctcaagcgc	ctccaccaag	ggcccatcgg	tcttcccc		408

<210> 72

<211> 414

<212> DNA

<213> Artificial Sequence

<220>

<223> Heavy Chain nucleic acid sequence

<400> 72

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tcttgcgctg	cttccggatt	cactttctct	ctttaccata	tggattgggt	tcgccaagct	120
cctggtaaag	gtttgagtg	ggtttctgtt	atctatcctt	ctggtggcgg	tactccttat	180
gctgactccg	ttaaagggtcg	cttcactatc	tctagagaca	actctaagaa	tactctctac	240
ttgcagatga	acagcttaag	ggctgaggac	actgcagtct	actattgtgc	gagacgggta	300
ggatattgta	gtggtggtag	ctgctactac	tactactact	acatggacgt	ctggggcaaa	360
gggaccacgg	tcaccgtctc	aagcgcctcc	accaagggcc	catcgttctt	cccg	414

<210> 73

<211> 396

<212> DNA

<213> Artificial Sequence

<220>

<223> Heavy Chain nucleic acid sequence

<400> 73

gaagttcaat	tgtttagagtc	tggtggcggt	cttggttcagc	ctggtgggttc	tttacgtctt	60
tcttgcgctg	cttccggatt	cactttctct	tggtactgga	tgaattgggt	tcgccaagct	120
cctggtaaag	gtttgagtg	ggtttcttct	atctattctt	ctggtggcta	tacttcttat	180
gctgactccg	ttaaagggtcg	cttcactatc	tctagagaca	actctaagaa	tactctctac	240
ttgcagatga	acagcttaag	ggctgaggac	actgcagtct	actattgtgc	gagagttcgg	300
gatattttga	ctggtcccta	ctactttgac	tactggggcc	agggaaccct	ggtcaccgtc	360
tcaagcgcct	ccaccaaggg	cccatcggtc	ttcccc			396

<210> 74

<211> 393
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Heavy Chain nucleic acid sequence

<400> 74
 gaagttcaat tgtagagtc tgggtggcggc cttgttcagc ctggtgggtc tttacgtctt 60
 tcttgcgctg cttccggatt cactttctct aattaccgta tgccttgggt tcgccaagct 120
 cctggtaaag gtttggagtg ggtttcttat atctattctt ctggtggcat tactcagtat 180
 gctgactccg ttaaagggtcg cttcactatc tctagagaca actctaagaa tactctctac 240
 ttgcagatga acagcttaag ggctgaggac actgcagtct actattgtgc gagatcgcca 300
 tcttactatg gttcggggtc gtcgcggtag tggggccagg gaaccctggc caccgtctca 360
 agcgctctca ccaagggccc atcgggtctc ccg 393

<210> 75
 <211> 405
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Heavy Chain nucleic acid sequence

<400> 75
 gaagttcaat tgtagagtc tgggtggcggc cttgttcagc ctggtgggtc tttacgtctt 60
 tcttgcgctg cttccggatt cactttctct cagtacatga tgacttgggt tcgccaagct 120
 cctggtaaag gtttggagtg ggtttcttat atcggttctt ctggtggcca gactaagtat 180
 gctgactccg ttaaagggtcg cttcactatc tctagagaca actctaagaa tactctctac 240
 ttgcagatga acagcttaag ggctgaggac actgcagtct actattgtgc gagggatcca 300
 ggggtagcag tggctgggta ctactactac ggtatggacg tctggggcca agggaccacg 360
 gtcaccgtct caagcgctc caccaagggc ccacgggtct tcccc 405

<210> 76
 <211> 411
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Heavy Chain nucleic acid sequence

<400> 76
 gaagttcaat tgtagagtc tgggtggcggc cttgttcagc ctggtgggtc tttacgtctt 60
 tcttgcgctg cttccggatt cactttctct cagtacaata tgccttgggt tcgccaagct 120
 cctggtaaag gtttggagtg ggtttcttct atcgttcctt ctggtgggtt tactgcttat 180
 gctgactccg ttaaagggtcg cttcactatc tctagagaca actctaagaa tactctctac 240
 ttgcagatga acagcttaag ggctgaggac actgcagtct actattgtgc gagagtcgat 300
 tgtagtgggt gtagctgcta ccgggggtccc caaaactact ttgactactg gggccagggg 360
 accctgggtc ccgtctcaag cgcctccacc aagggcccat cggctcttccc g 411

<210> 77
 <211> 351
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Heavy Chain nucleic acid sequence

<400> 77

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gaagttcaat tgtagagtc tggtagcggt cttgttcagc ctggtgggtc tttacgtctt      60
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cctggtaagg tttggagtgg gtttctgtta tcgtttcttc tggtagcact actgagtatg      180
ctgactccgt taaaggtcgc ttcactatct ctagagacaa ctctaagaat actctctact      240
tgcagatgaa cagcttaagg gctgaggaca ctgcagtcta ctattgtgcg agagggggat      300
attgtagtgg tggcaggtgt tacacctggc tcgaagacta ctggggccag g              351

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<210> 78

<211> 110

<212> PRT

<213> Artificial Sequence

<220>

<223> Light Chain amino acid sequence

<400> 78

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Gln Ser Val Leu Thr Gln Pro Pro Ser Ala Ser Gly Thr Pro Gly Gln
 1           5           10           15
Arg Val Thr Ile Ser Cys Ser Gly Ser Ser Ser Asn Ile Glu Ser Asn
      20           25           30
Thr Val Thr Trp Tyr Gln Gln Leu Pro Gly Thr Ala Pro Lys Leu Leu
      35           40           45
Ile Tyr Ser Asp Asp Gln Arg Pro Ser Gly Val Pro Asp Arg Phe Ser
      50           55           60
Gly Ser Lys Ser Gly Thr Ser Ala Ser Leu Ala Ile Ser Gly Leu Gln
65           70           75           80
Ser Glu Asp Glu Ala Asp Tyr Tyr Cys Ala Thr Trp Asp Asn Thr Leu
      85           90           95
Arg Gly Val Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
      100           105           110

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<210> 79

<211> 132

<212> PRT

<213> Artificial Sequence

<220>

<223> Heavy Chain amino acid sequence

<400> 79

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Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
 1           5           10           15
Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Pro Tyr
      20           25           30
Arg Met Asp Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
      35           40           45
Ser Tyr Ile Tyr Pro Ser Gly Gly Phe Thr Pro Tyr Ala Asp Ser Val
      50           55           60
Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Phe Tyr
65           70           75           80
Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
      85           90           95
Ala Lys Gly Ser Thr Gly Tyr Arg Tyr Tyr Tyr Gly Met Asp Val Trp
      100           105           110
Gly Gln Gly Thr Thr Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro

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115 120 125
 Ser Val Phe Pro
 130

<210> 80
 <211> 13
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Synthetically generated peptide

<400> 80
 Ser Gly Ser Ser Ser Asn Ile Glu Ser Asn Thr Val Thr
 1 5 10

<210> 81
 <211> 7
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Synthetically generated peptide

<400> 81
 Ser Asp Asp Gln Arg Pro Ser
 1 5

<210> 82
 <211> 11
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Synthetically generated peptide

<400> 82
 Ala Thr Trp Asp Asn Thr Leu Arg Gly Val Val
 1 5 10

<210> 83
 <211> 5
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Synthetically generated peptide

<400> 83
 Pro Tyr Arg Met Asp
 1 5

<210> 84
 <211> 17
 <212> PRT
 <213> Artificial Sequence

<220>

<223> Synthetically generated peptide

<400> 84

Tyr Ile Tyr Pro Ser Gly Gly Phe Thr Pro Tyr Ala Asp Ser Val Lys
 1 5 10 15
 Gly

<210> 85

<211> 13

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetically generated peptide

<400> 85

Gly Ser Thr Gly Tyr Arg Tyr Tyr Tyr Gly Met Asp Val
 1 5 10

<210> 86

<211> 120

<212> PRT

<213> Artificial Sequence

<220>

<223> Light Chain amino acid sequence

<400> 86

Gln Asp Ile Val Met Thr Gln Thr Pro Pro Ser Leu Pro Val Asn Pro
 1 5 10 15
 Gly Glu Pro Ala Ser Ile Ser Cys Lys Ser Ser Gln Ser Leu Leu Gln
 20 25 30
 Ser Asn Gly Tyr Asn Tyr Leu Asp Trp Tyr Leu Gln Lys Pro Gly Gln
 35 40 45
 Ser Pro Gln Leu Leu Ile Tyr Leu Gly Ser Asn Arg Ala Ser Gly Val
 50 55 60
 Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Lys
 65 70 75 80
 Ile Ser Arg Val Glu Ala Glu Asp Val Gly Ile Tyr Tyr Cys Met Gln
 85 90 95
 Ala Leu His Thr Pro Pro Phe Gly Gln Gly Thr Arg Leu Glu Ile Lys
 100 105 110
 Arg Thr Val Ala Ala Pro Ser Val
 115 120

<210> 87

<211> 132

<212> PRT

<213> Artificial Sequence

<220>

<223> Heavy Chain amino acid sequence

<400> 87

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly

1				5					10					15			
Ser	Leu	Arg	Leu	Ser	Cys	Ala	Ala	Ser	Gly	Phe	Thr	Phe	Ser	Trp	Tyr		
			20					25					30				
Trp	Met	Asn	Trp	Val	Arg	Gln	Ala	Pro	Gly	Lys	Gly	Leu	Glu	Trp	Val		
		35					40					45					
Ser	Ser	Ile	Tyr	Ser	Ser	Gly	Gly	Tyr	Thr	Ser	Tyr	Ala	Asp	Ser	Val		
	50					55					60						
Lys	Gly	Arg	Phe	Thr	Ile	Ser	Arg	Asp	Asn	Ser	Lys	Asn	Thr	Leu	Tyr		
65					70				75					80			
Leu	Gln	Met	Asn	Ser	Leu	Arg	Ala	Glu	Asp	Thr	Ala	Val	Tyr	Tyr	Cys		
			85						90				95				
Ala	Arg	Val	Arg	Asp	Ile	Leu	Thr	Gly	Pro	Tyr	Tyr	Phe	Asp	Tyr	Trp		
			100					105					110				
Gly	Gln	Gly	Thr	Leu	Val	Thr	Val	Ser	Ser	Ala	Ser	Thr	Lys	Gly	Pro		
		115					120					125					
Ser	Val	Phe	Pro														
	130																

<210> 88

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetically generated peptide

<400> 88

Lys	Ser	Ser	Gln	Ser	Leu	Leu	Gln	Ser	Asn	Gly	Tyr	Asn	Tyr	Leu	Asp
1				5					10					15	

<210> 89

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetically generated peptide

<400> 89

Leu	Gly	Ser	Asn	Arg	Ala	Ser
1				5		

<210> 90

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetically generated peptide

<400> 90

Met	Gln	Ala	Leu	His	Thr	Pro	Pro
1				5			

<210> 91

<211> 5

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetically generated peptide

<400> 91

Trp Tyr Trp Met Asn
1 5

<210> 92

<211> 17

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetically generated peptide

<400> 92

Ser Ile Tyr Ser Ser Gly Gly Tyr Thr Ser Tyr Ala Asp Ser Val Lys
1 5 10 15
Gly

<210> 93

<211> 13

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetically generated peptide

<400> 93

Val Arg Asp Ile Leu Thr Gly Pro Tyr Tyr Phe Asp Tyr
1 5 10

<210> 94

<211> 116

<212> PRT

<213> Artificial Sequence

<220>

<223> Light Chain amino acid sequence

<400> 94

Gln Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val
1 5 10 15
Gly Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Gly Ile Arg His
20 25 30
Tyr Leu Gly Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Arg Leu
35 40 45
Ile Tyr Ala Ala Ser Ser Leu Gln Phe Gly Val Pro Ala Arg Phe Ser
50 55 60
Gly Ser Gly Ser Gly Thr Glu Phe Thr Leu Thr Ile Ser Ser Leu Gln
65 70 75 80
Pro Glu Asp Phe Ala Thr Tyr Tyr Cys Leu Gln His Asn Ser Phe Pro
85 90 95
Pro Ala Phe Gly Gln Gly Thr Lys Val Glu Ile Lys Arg Thr Val Ala

100 105 110
 Ala Pro Ser Val
 115

<210> 95
 <211> 132
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 95
 Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
 1 5 10 15
 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Pro Tyr
 20 25 30
 Asp Met Trp Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
 35 40 45
 Ser Tyr Ile Ser Ser Ser Gly Gly Lys Thr Met Tyr Ala Asp Ser Val
 50 55 60
 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
 65 70 75 80
 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
 85 90 95
 Ala Arg Leu Gly Gly Asn Ser His Tyr Tyr Tyr Gly Met Asp Val Trp
 100 105 110
 Gly Gln Gly Thr Thr Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro
 115 120 125
 Ser Val Phe Pro
 130

<210> 96
 <211> 11
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Synthetically generated peptide

<400> 96
 Arg Ala Ser Gln Gly Ile Arg His Tyr Leu Gly
 1 5 10

<210> 97
 <211> 7
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Synthetically generated peptide

<400> 97
 Ala Ala Ser Ser Leu Gln Phe
 1 5

<210> 98

<211> 9
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Synthetically generated peptide

<400> 98
 Leu Gln His Asn Ser Phe Pro Pro Ala
 1 5

<210> 99
 <211> 5
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Synthetically generated peptide

<400> 99
 Pro Tyr Asp Met Trp
 1 5

<210> 100
 <211> 17
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Synthetically generated peptide

<400> 100
 Tyr Ile Ser Ser Ser Gly Gly Lys Thr Met Tyr Ala Asp Ser Val Lys
 1 5 10 15
 Gly

<210> 101
 <211> 13
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Synthetically generated peptide

<400> 101
 Leu Gly Gly Asn Ser His Tyr Tyr Tyr Gly Met Asp Val
 1 5 10

<210> 102
 <211> 116
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 102

```

Gln Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Val Ser Ala Ser Val
 1           5           10           15
Gly Asp Arg Ile Ala Ile Thr Cys Arg Ala Ser Gln Gly Ile Ser Thr
          20           25           30
Trp Leu Ala Trp Tyr Gln Gln Arg Pro Gly Arg Ala Pro Lys Leu Leu
          35           40           45
Ile Tyr Ala Ala Ser Thr Leu Gln Ser Gly Val Pro Ser Arg Phe Ser
          50           55           60
Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln
65          70          75          80
Pro Glu Asp Phe Ala Thr Tyr Phe Cys Gln Gln Ala Asp Ser Phe Pro
          85          90          95
Leu Thr Phe Gly Gln Gly Thr Lys Leu Glu Ile Lys Arg Thr Val Ala
          100         105         110
Ala Pro Ser Val
          115

```

<210> 103

<211> 123

<212> PRT

<213> Artificial Sequence

<220>

<223> Heavy Chain amino acid sequence

Heavy Chain amino acid sequence

Heavy Chain amino acid sequence

Heavy Chain amino acid sequence

Heavy Chain amino acid sequence

<400> 103

```

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
 1           5           10           15
Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Asn Tyr
          20           25           30
Ala Met Asp Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
          35           40           45
Ser Tyr Ile Ser Pro Ser Gly Gly Tyr Thr Arg Tyr Ala Asp Ser Val
          50           55           60
Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65          70          75          80
Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
          85          90          95
Ala Ser Asp Phe Gly Ser Trp Gly Gln Gly Thr Leu Val Thr Val Ser
          100         105         110
Ser Ala Ser Thr Lys Gly Pro Ser Val Phe Pro
          115         120

```

<210> 104

<211> 11

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetically generated peptide

<400> 104

Arg Ala Ser Gln Gly Ile Ser Thr Trp Leu Ala
1 5 10

<210> 105

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetically generated peptide

<400> 105

Ala Ala Ser Thr Leu Gln Ser
1 5

<210> 106

<211> 9

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetically generated peptide

<400> 106

Gln Gln Ala Asp Ser Phe Pro Leu Thr
1 5

<210> 107

<211> 5

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetically generated peptide

<400> 107

Asn Tyr Ala Met Asp
1 5

<210> 108

<211> 17

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetically generated peptide

<400> 108

Tyr Ile Ser Pro Ser Gly Gly Tyr Thr Arg Tyr Ala Asp Ser Val Lys
1 5 10 15
Gly

<210> 109
 <211> 4
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Synthetically generated peptide

<400> 109
 Asp Phe Gly Ser
 1

<210> 110
 <211> 117
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 110
 Gln Asp Ile Gln Met Thr Gln Ser Pro Gly Thr Leu Ser Leu Ser Pro
 1 5 10 15
 Gly Glu Arg Ala Thr Leu Ser Cys Arg Ala Ser Gln Ser Ile Ser Ser
 20 25 30
 Ser Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro Arg Leu
 35 40 45
 Leu Ile Tyr Ala Ala Ala Ser Arg Ala Thr Gly Ile Pro Asp Arg Phe
 50 55 60
 Ser Gly Ile Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu
 65 70 75 80
 Glu Pro Glu Asp Phe Ala Val Tyr Tyr Cys Gln Gln Arg Ser Asn Trp
 85 90 95
 Pro Leu Thr Phe Gly Gly Gly Thr Lys Val Glu Ile Lys Arg Thr Val
 100 105 110
 Ala Ala Pro Ser Val
 115

<210> 111
 <211> 131
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 111
 Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
 1 5 10 15
 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Arg Tyr
 20 25 30
 His Met Glu Trp Val Arg Gln Ala His Gly Lys Gly Leu Glu Trp Val
 35 40 45
 Ser Tyr Ile Ser Pro Ser Gly Gly Lys Thr Leu Tyr Ala Asp Ser Val
 50 55 60
 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
 65 70 75 80

Leu	Gln	Met	Asn	Ser	Leu	Arg	Ala	Glu	Asp	Thr	Ala	Val	Tyr	Tyr	Cys
				85					90					95	
Ala	Arg	His	Leu	Gly	Tyr	Gly	Ser	Gly	Ser	Tyr	Phe	Asp	Tyr	Trp	Gly
			100					105					110		
Gln	Gly	Thr	Leu	Val	Thr	Val	Ser	Ser	Ala	Ser	Thr	Lys	Gly	Pro	Ser
		115					120						125		
Val	Phe	Pro													
		130													

<210> 112
 <211> 12
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Synthetically generated peptide

<400>	112										
Arg	Ala	Ser	Gln	Ser	Ile	Ser	Ser	Ser	Tyr	Leu	Ala
1				5					10		

<210> 113
 <211> 7
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Synthetically generated peptide

<400>	113					
Ala	Ala	Ala	Ser	Arg	Ala	Thr
1				5		

<210> 114
 <211> 9
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Synthetically generated peptide

<400>	114							
Gln	Gln	Arg	Ser	Asn	Trp	Pro	Leu	Thr
1				5				

<210> 115
 <211> 5
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Synthetically generated peptide

<400>	115			
Arg	Tyr	His	Met	Glu
1			5	

<210> 116
 <211> 17
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Synthetically generated peptide

<400> 116
 Tyr Ile Ser Pro Ser Gly Gly Lys Thr Leu Tyr Ala Asp Ser Val Lys
 1 5 10 15
 Gly

<210> 117
 <211> 12
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Synthetically generated peptide

<400> 117
 His Leu Gly Tyr Gly Ser Gly Ser Tyr Phe Asp Tyr
 1 5 10

<210> 118
 <211> 116
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 118
 Gln Tyr Glu Leu Thr Gln Pro Pro Ser Val Ser Val Ser Pro Gly Gln
 1 5 10 15
 Thr Ala Thr Ile Ile Cys Ser Gly Asp Lys Leu Gly Asp Lys Tyr Val
 20 25 30
 Ala Trp Tyr Gln Gln Lys Pro Gly Gln Ser Pro Val Leu Val Val Tyr
 35 40 45
 Glu Asp Asn Lys Arg Pro Ser Gly Ile Pro Glu Arg Ile Ser Gly Ser
 50 55 60
 Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
 65 70 75 80
 Asp Asp Ala Asp Tyr Tyr Cys Gln Ala Trp Asp Arg Ser Thr Asp His
 85 90 95
 Tyr Val Phe Gly Thr Gly Thr Lys Val Thr Val Leu Gly Gln Pro Lys
 100 105 110
 Ala Asn Pro Thr
 115

<210> 119
 <211> 131
 <212> PRT
 <213> Artificial Sequence

<220>

<223> Heavy Chain amino acid sequence

<400> 119

```

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
 1             5             10             15
Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Asn Tyr
          20             25             30
Arg Met Pro Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
          35             40             45
Ser Tyr Ile Tyr Ser Ser Gly Gly Ile Thr Gln Tyr Ala Asp Ser Val
 50             55             60
Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65             70             75             80
Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
          85             90             95
Ala Arg Ser Arg Ser Tyr Tyr Gly Ser Gly Ser Ser Arg Tyr Trp Gly
          100             105             110
Gln Gly Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser
          115             120             125
Val Phe Pro
          130

```

<210> 120

<211> 11

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetically generated peptide

<400> 120

```

Ser Gly Asp Lys Leu Gly Asp Lys Tyr Val Ala
 1             5             10

```

<210> 121

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetically generated peptide

<400> 121

```

Glu Asp Asn Lys Arg Pro Ser
 1             5

```

<210> 122

<211> 11

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetically generated peptide

<400> 122

```

Gln Ala Trp Asp Arg Ser Thr Asp His Tyr Val

```

1 5 10

<210> 123

<211> 5

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetically generated peptide

<400> 123

Asn Tyr Arg Met Pro

1 5

<210> 124

<211> 17

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetically generated peptide

<400> 124

Tyr Ile Tyr Ser Ser Gly Gly Ile Thr Gln Tyr Ala Asp Ser Val Lys

1 5 10 15

Gly

<210> 125

<211> 12

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetically generated peptide

<400> 125

Ser Arg Ser Tyr Tyr Gly Ser Gly Ser Ser Arg Tyr

1 5 10

<210> 126

<211> 108

<212> PRT

<213> Unknown

<220>

<223> Synthetically generated peptide

<223> Light Chain amino acid sequence

<400> 126

Gln Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Phe Ser Ala Ser Thr

1 5 10 15

Gly Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Gly Ile Ser Ser

20 25 30

Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu

35 40 45

```

Ile Tyr Ala Ala Ser Thr Leu Gln Ser Gly Val Pro Ser Lys Phe Ser
 50                      55                      60
Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln
65                      70                      75                      80
Pro Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Tyr Asn Ser Tyr Pro
                      85                      90                      95
Leu Thr Phe Gly Gln Gly Thr Arg Leu Glu Ile Lys
                100                      105

```

```

<210> 127
<211> 123
<212> PRT
<213> Artificial Sequence

```

```

<220>
<223> Heavy Chain amino acid sequence

```

```

<400> 127
Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
 1                      5                      10                      15
Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Trp Tyr
                20                      25                      30
Thr Met Val Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
                35                      40                      45
Ser Ser Ile Tyr Ser Ser Gly Gly Phe Thr Trp Tyr Ala Asp Ser Val
 50                      55                      60
Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65                      70                      75                      80
Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
                85                      90                      95
Ala Ser Asp Phe Gly Ser Trp Gly Gln Gly Thr Leu Val Thr Val Ser
                100                      105                      110
Ser Ala Ser Thr Lys Gly Pro Ser Val Phe Pro
                115                      120

```

```

<210> 128
<211> 11
<212> PRT
<213> Artificial Sequence

```

```

<220>
<223> Synthetically generated peptide

```

```

<400> 128
Arg Ala Ser Gln Gly Ile Ser Ser Tyr Leu Ala
 1                      5                      10

```

```

<210> 129
<211> 7
<212> PRT
<213> Artificial Sequence

```

```

<220>
<223> Synthetically generated peptide

```

```

<400> 129
Ala Ala Ser Thr Leu Gln Ser

```

```

1              5

<210> 130
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetically generated peptide

<400> 130
Gln Gln Tyr Asn Ser Tyr Pro Leu Thr
1              5

<210> 131
<211> 5
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetically generated peptide

<400> 131
Trp Tyr Thr Met Val
1              5

<210> 132
<211> 17
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetically generated peptide

<400> 132
Ser Ile Tyr Ser Ser Gly Gly Phe Thr Trp Tyr Ala Asp Ser Val Lys
1              5              10              15
Gly

<210> 133
<211> 4
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetically generated peptide

<400> 133
Asp Phe Gly Ser
1

<210> 134
<211> 116
<212> PRT
<213> Artificial Sequence

```

<220>

<223> Light Chain amino acid sequence

<400> 134

Gln	Asp	Ile	Gln	Met	Thr	Gln	Ser	Pro	Ser	Ser	Leu	Tyr	Ala	Ser	Val
1				5				10						15	
Gly	Asp	Arg	Val	Thr	Ile	Thr	Cys	Arg	Ala	Ser	Gln	Gly	Ile	Arg	Asn
			20					25					30		
Glu	Leu	Gly	Trp	Tyr	Gln	Gln	Lys	Pro	Gly	Lys	Ala	Pro	Gln	Arg	Leu
		35					40					45			
Ile	Tyr	Asp	Ala	Ser	Thr	Leu	Gln	Ser	Gly	Val	Pro	Ser	Arg	Phe	Ser
	50					55					60				
Gly	Gly	Gly	Ser	Arg	Thr	Glu	Phe	Thr	Leu	Thr	Ile	Ser	Ser	Leu	Glu
65					70					75				80	
Pro	His	Asp	Phe	Gly	Thr	Tyr	Tyr	Cys	Gln	Gln	Tyr	Ala	Ser	Tyr	Pro
				85					90					95	
Leu	Thr	Phe	Gly	Gly	Gly	Thr	Lys	Val	Glu	Ile	Lys	Arg	Thr	Val	Ala
			100					105						110	
Ala	Pro	Ser	Val												
			115												

<210> 135

<211> 140

<212> PRT

<213> Artificial Sequence

<220>

<223> Heavy Chain amino acid sequence

<400> 135

Glu	Val	Gln	Leu	Leu	Glu	Ser	Gly	Gly	Gly	Leu	Val	Gln	Pro	Gly	Gly
1				5					10					15	
Ser	Leu	Arg	Leu	Ser	Cys	Ala	Ala	Ser	Gly	Phe	Thr	Phe	Ser	Asp	Tyr
			20					25					30		
Lys	Met	Pro	Trp	Val	Arg	Gln	Ala	Pro	Gly	Lys	Gly	Leu	Glu	Trp	Val
		35					40					45			
Ser	Ser	Ile	Trp	Ser	Ser	Gly	Gly	Thr	Thr	Glu	Tyr	Ala	Asp	Ser	Val
	50					55					60				
Lys	Gly	Arg	Phe	Thr	Ile	Ser	Arg	Asp	Asn	Ser	Lys	Asn	Thr	Leu	Tyr
65					70					75				80	
Leu	Gln	Met	Asn	Ser	Leu	Arg	Ala	Glu	Asp	Thr	Ala	Val	Tyr	Tyr	Cys
			85						90					95	
Ala	Arg	Glu	Glu	Ile	Gly	Arg	Tyr	Phe	Asp	Trp	Phe	Leu	Gly	Asn	Tyr
			100					105					110		
Tyr	Tyr	Tyr	Gly	Met	Asp	Val	Trp	Gly	Gln	Gly	Thr	Thr	Val	Thr	Val
		115				120						125			
Ser	Ser	Ala	Ser	Thr	Lys	Gly	Pro	Ser	Val	Phe	Pro				
		130				135					140				

<210> 136

<211> 11

<212> PRT

<213> Artificial Sequence

<220>

<223> Light Chain amino acid sequence

<400> 136
 Arg Ala Ser Gln Gly Ile Arg Asn Glu Leu Gly
 1 5 10

<210> 137
 <211> 7
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 137
 Asp Ala Ser Thr Leu Gln Ser
 1 5

<210> 138
 <211> 9
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 138
 Gln Gln Tyr Ala Ser Tyr Pro Leu Thr
 1 5

<210> 139
 <211> 5
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 139
 Asp Tyr Lys Met Pro
 1 5

<210> 140
 <211> 17
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 140
 Ser Ile Trp Ser Ser Gly Gly Thr Thr Glu Tyr Ala Asp Ser Val Lys
 1 5 10 15
 Gly

<210> 141
 <211> 21
 <212> PRT

<213> Artificial Sequence

<220>

<223> Heavy Chain amino acid sequence

<400> 141

Glu	Glu	Ile	Gly	Arg	Tyr	Phe	Asp	Trp	Phe	Leu	Gly	Asn	Tyr	Tyr	Tyr
1				5					10					15	
Tyr	Gly	Met	Asp	Val											
				20											

<210> 142

<211> 118

<212> PRT

<213> Artificial Sequence

<220>

<223> Light Chain amino acid sequence

<400> 142

Gln	Ser	Ala	Leu	Thr	Gln	Pro	Pro	Ser	Ala	Ser	Gly	Thr	Pro	Gly	Gln
1				5					10					15	
Arg	Val	Thr	Ile	Ser	Cys	Ser	Gly	Ser	Ser	Ser	Asn	Ile	Gly	Ser	Asn
			20				25					30			
Phe	Val	Tyr	Trp	Tyr	His	His	Leu	Pro	Gly	Thr	Ala	Pro	Lys	Leu	Leu
		35					40				45				
Ile	Tyr	Arg	Asn	Asn	Gln	Arg	Pro	Ser	Gly	Val	Pro	Asp	Arg	Phe	Ser
	50				55					60					
Gly	Ser	Lys	Ser	Gly	Thr	Ser	Ala	Ser	Leu	Ala	Ile	Ser	Gly	Leu	Arg
65				70					75					80	
Ser	Glu	Asp	Glu	Ala	Asp	Tyr	Tyr	Cys	Ala	Ala	Trp	Asp	Asp	Ser	Leu
			85					90						95	
Ser	Gly	Val	Val	Phe	Gly	Gly	Gly	Thr	Lys	Leu	Thr	Val	Leu	Gly	Gln
		100					105					110			
Pro	Lys	Ala	Ala	Pro	Ser										
			115												

<210> 143

<211> 128

<212> PRT

<213> Artificial Sequence

<220>

<223> Heavy Chain amino acid sequence

<400> 143

Glu	Val	Gln	Leu	Leu	Glu	Ser	Gly	Gly	Gly	Leu	Val	Gln	Pro	Gly	Gly
1				5					10					15	
Ser	Leu	Arg	Leu	Ser	Cys	Ala	Ala	Ser	Gly	Phe	Thr	Phe	Ser	Gln	Tyr
		20					25					30			
Lys	Met	Asn	Trp	Val	Arg	Gln	Ala	Pro	Gly	Lys	Gly	Leu	Glu	Trp	Val
		35				40					45				
Ser	Tyr	Ile	Ser	Pro	Ser	Gly	Gly	Tyr	Thr	Ala	Tyr	Ala	Asp	Ser	Val
	50				55					60					
Lys	Gly	Arg	Phe	Thr	Ile	Ser	Arg	Asp	Asn	Ser	Lys	Asn	Thr	Leu	Tyr
65				70					75					80	
Leu	Gln	Met	Asn	Ser	Leu	Arg	Ala	Glu	Asp	Thr	Ala	Val	Tyr	Tyr	Cys

				85					90					95					
Ala	Arg	Asp	Val	Val	Ala	Gly	Pro	Phe	Asp	Tyr	Trp	Gly	Gln	Gly	Thr				
			100					105					110						
Leu	Val	Thr	Val	Ser	Ser	Ala	Ser	Thr	Lys	Gly	Pro	Ser	Val	Phe	Pro				
		115					120					125							

<210> 144
 <211> 13
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 144
 Ser Gly Ser Ser Ser Asn Ile Gly Ser Asn Phe Val Tyr
 1 5 10

<210> 145
 <211> 7
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 145
 Arg Asn Asn Gln Arg Pro Ser
 1 5

<210> 146
 <211> 11
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 146
 Ala Ala Trp Asp Asp Ser Leu Ser Gly Val Val
 1 5 10

<210> 147
 <211> 5
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 147
 Gln Tyr Lys Met Asn
 1 5

<210> 148
 <211> 17
 <212> PRT

<213> Artificial Sequence

<220>

<223> Heavy Chain amino acid sequence

<400> 148

Tyr Ile Ser Pro Ser Gly Gly Tyr Thr Ala Tyr Ala Asp Ser Val Lys
 1 5 10 15
 Gly

<210> 149

<211> 9

<212> PRT

<213> Artificial Sequence

<220>

<223> Heavy Chain amino acid sequence

<400> 149

Asp Val Val Ala Gly Pro Phe Asp Tyr
 1 5

<210> 150

<211> 116

<212> PRT

<213> Artificial Sequence

<220>

<223> Light Chain amino acid sequence

<400> 150

Gln Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val
 1 5 10 15
 Gly Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Asp Ile Ser Asn
 20 25 30
 Tyr Leu Ala Trp Phe Gln Gln Lys Pro Gly Arg Ala Pro Lys Ser Leu
 35 40 45
 Ile Tyr Gly Ala Ser Ser Leu Gln Thr Gly Val Pro Ser Lys Phe Ser
 50 55 60
 Gly Ser Gly Ser Gly Thr Glu Phe Thr Leu Thr Ile Ser Gly Leu Gln
 65 70 75 80
 Pro Glu Asp Val Ala Thr Tyr Tyr Cys His Gln Tyr Asn His Tyr Pro
 85 90 95
 Pro Thr Phe Gly Gly Gly Thr Lys Val Glu Ile Lys Arg Thr Val Ala
 100 105 110
 Ala Pro Ser Val
 115

<210> 151

<211> 129

<212> PRT

<213> Artificial Sequence

<220>

<223> Heavy Chain amino acid sequence

<400> 151
 Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
 1 5 10 15
 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Lys Tyr
 20 25 30
 Pro Met Phe Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
 35 40 45
 Ser Trp Ile Ser Pro Ser Gly Gly Lys Thr Val Tyr Ala Asp Ser Val
 50 55 60
 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
 65 70 75 80
 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
 85 90 95
 Ala Lys Asp Cys Arg Gly Gly Cys Ser Gly Gly Ser Trp Gly Gln Gly
 100 105 110
 Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val Phe
 115 120 125
 Pro

<210> 152
 <211> 11
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 152
 Arg Ala Ser Gln Asp Ile Ser Asn Tyr Leu Ala
 1 5 10

<210> 153
 <211> 7
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 153
 Gly Ala Ser Ser Leu Gln Thr
 1 5

<210> 154
 <211> 9
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 154
 His Gln Tyr Asn His Tyr Pro Pro Thr
 1 5

<210> 155

<211> 5
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 155
 Lys Tyr Pro Met Phe
 1 5

<210> 156
 <211> 17
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 156
 Trp Ile Ser Pro Ser Gly Gly Lys Thr Val Tyr Ala Asp Ser Val Lys
 1 5 10 15
 Gly

<210> 157
 <211> 10
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 157
 Asp Cys Arg Gly Gly Cys Ser Gly Gly Ser
 1 5 10

<210> 158
 <211> 116
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 158
 Gln Asp Ile Gln Met Thr Gln Ser Pro Ala Thr Leu Ser Val Ser Pro
 1 5 10 15
 Gly Glu Arg Ala Thr Leu Ser Cys Arg Ala Ser Gln Asp Val Asn Arg
 20 25 30
 Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Gln Pro Pro Arg Leu Leu
 35 40 45
 Ile Tyr Gly Ala Ser Thr Arg Ala Thr Gly Ile Pro Ala Arg Ile Ser
 50 55 60
 Gly Ser Gly Ser Gly Thr Glu Phe Thr Leu Thr Ile Ser Ser Leu Gln
 65 70 75 80
 Ser Glu Asp Phe Ala Val Tyr Tyr Cys Gln Gln Tyr His Asn Trp Pro

```

      85          90          95
Leu Thr Phe Gly Gly Gly Thr Lys Val Glu Ile Lys Arg Thr Val Ala
      100          105          110
Ala Pro Ser Val
      115

```

```

<210> 159
<211> 123
<212> PRT
<213> Artificial Sequence

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```

<220>
<223> Heavy Chain amino acid sequence

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```

<400> 159
Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
 1          5          10          15
Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Arg Tyr
      20          25          30
Ser Met Asn Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
      35          40          45
Ser Tyr Ile Ser Pro Ser Gly Gly Met Thr Lys Tyr Ala Asp Ser Val
      50          55          60
Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
      65          70          75          80
Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
      85          90          95
Ala Asn Thr Leu Gly Tyr Trp Gly Gln Gly Thr Leu Val Thr Val Ser
      100          105          110
Ser Ala Ser Thr Lys Gly Pro Ser Val Phe Pro
      115          120

```

```

<210> 160
<211> 11
<212> PRT
<213> Artificial Sequence

```

```

<220>
<223> Light Chain amino acid sequence

```

```

<400> 160
Arg Ala Ser Gln Asp Val Asn Arg Tyr Leu Ala
 1          5          10

```

```

<210> 161
<211> 7
<212> PRT
<213> Artificial Sequence

```

```

<220>
<223> Light Chain amino acid sequence

```

```

<400> 161
Gly Ala Ser Thr Arg Ala Thr
 1          5

```

```

<210> 162

```

<211> 9
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 162
 Gln Gln Tyr His Asn Trp Pro Leu Thr
 1 5

<210> 163
 <211> 5
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 163
 Arg Tyr Ser Met Asn
 1 5

<210> 164
 <211> 17
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 164
 Tyr Ile Ser Pro Ser Gly Gly Met Thr Lys Tyr Ala Asp Ser Val Lys
 1 5 10 15
 Gly

<210> 165
 <211> 4
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 165
 Thr Leu Gly Tyr
 1

<210> 166
 <211> 119
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 166

```

Gln Ser Ala Leu Thr Gln Pro Ala Ser Val Ser Gly Ser Pro Gly Gln
 1           5           10           15
Ser Ile Thr Ile Ser Cys Thr Gly Thr Ser Ser Asp Val Gly Tyr Tyr
          20           25           30
Asp Tyr Val Ser Trp Tyr Gln His His Pro Gly Lys Ala Pro Lys Leu
      35           40           45
Ile Ile Tyr Asp Val Thr Ser Arg Pro Ser Gly Val Ser Ser His Phe
 50           55           60
Ser Gly Ser Lys Ser Gly Asn Thr Ala Ser Leu Thr Ile Ser Gly Leu
65           70           75           80
Gln Ala Asp Asp Glu Ala Asp Tyr Tyr Cys Ser Ser Tyr Thr Ser Gly
          85           90           95
Ser Thr Arg Tyr Val Phe Gly Pro Gly Thr Lys Val Thr Val Leu Gly
          100          105          110
Gln Pro Lys Ala Asn Pro Thr
          115

```

<210> 167

<211> 131

<212> PRT

<213> Artificial Sequence

<220>

<223> Heavy Chain amino acid sequence

<400> 167

```

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
 1           5           10           15
Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Asp Tyr
          20           25           30
Tyr Met Arg Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
      35           40           45
Ser Arg Ile Tyr Pro Ser Gly Gly His Thr Trp Tyr Ala Asp Ser Val
 50           55           60
Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65           70           75           80
Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
          85           90           95
Ala Arg His Arg Ala Gly Ser Ser Gly Trp Tyr Ser Asp Tyr Trp Gly
          100          105          110
Gln Gly Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser
          115          120          125
Val Phe Pro
          130

```

<210> 168

<211> 14

<212> PRT

<213> Artificial Sequence

<220>

<223> Light Chain amino acid sequence

<400> 168

```

Thr Gly Thr Ser Ser Asp Val Gly Tyr Tyr Asp Tyr Val Ser
 1           5           10

```


<210> 169
 <211> 7
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 169
 Asp Val Thr Ser Arg Pro Ser
 1 5

<210> 170
 <211> 11
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 170
 Ser Ser Tyr Thr Ser Gly Ser Thr Arg Tyr Val
 1 5 10

<210> 171
 <211> 5
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 171
 Asp Tyr Tyr Met Arg
 1 5

<210> 172
 <211> 17
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 172
 Arg Ile Tyr Pro Ser Gly Gly His Thr Trp Tyr Ala Asp Ser Val Lys
 1 5 10 15
 Gly

<210> 173
 <211> 12
 <212> PRT
 <213> Artificial Sequence

<220>

<223> Heavy Chain amino acid sequence

<400> 173

His Arg Ala Gly Ser Ser Gly Trp Tyr Ser Asp Tyr
1 5 10

<210> 174

<211> 116

<212> PRT

<213> Artificial Sequence

<220>

<223> Light Chain amino acid sequence

<400> 174

Gln Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val
1 5 10 15
Gly Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Asp Ile Arg Asn
20 25 30
Tyr Leu Ala Trp Phe Gln Gln Lys Pro Gly Glu Ala Pro Lys Ser Leu
35 40 45
Ile Tyr Ala Ala Ser Ser Leu Gln Ser Gly Val Ser Ser Asn Phe Ser
50 55 60
Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln
65 70 75 80
Pro Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Tyr His Arg Tyr Pro
85 90 95
Arg Thr Phe Gly Gln Gly Thr Lys Leu Glu Ile Lys Arg Thr Val Ala
100 105 110
Ala Pro Ser Val
115

<210> 175

<211> 131

<212> PRT

<213> Artificial Sequence

<220>

<223> Heavy Chain amino acid sequence

<400> 175

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1 5 10 15
Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ala Tyr
20 25 30
Asn Met Pro Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45
Ser Tyr Ile Ser Ser Ser Gly Thr Gly Tyr Ala Asp Ser Val Lys Gly
50 55 60
Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr Leu Gln
65 70 75 80
Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys Ala Arg
85 90 95
Glu Leu Gly Ser Gly Ser Tyr Tyr Pro Gly Tyr Phe Gln His Trp Gly
100 105 110
Gln Gly Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser
115 120 125

Val Phe Pro
130

<210> 176
<211> 11
<212> PRT
<213> Artificial Sequence

<220>
<223> Light Chain amino acid sequence

<400> 176
Arg Ala Ser Gln Asp Ile Arg Asn Tyr Leu Ala
1 5 10

<210> 177
<211> 7
<212> PRT
<213> Artificial Sequence

<220>
<223> Light Chain amino acid sequence

<400> 177
Ala Ala Ser Ser Leu Gln Ser
1 5

<210> 178
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Light Chain amino acid sequence

<400> 178
Gln Gln Tyr His Arg Tyr Pro Arg Thr
1 5

<210> 179
<211> 5
<212> PRT
<213> Artificial Sequence

<220>
<223> Heavy Chain amino acid sequence

<400> 179
Ala Tyr Asn Met Pro
1 5

<210> 180
<211> 16
<212> PRT
<213> Artificial Sequence

<220>

<223> Heavy Chain amino acid sequence

<400> 180

Tyr Ile Ser Ser Ser Gly Thr Gly Tyr Ala Asp Ser Val Lys Gly Arg
1 5 10 15

<210> 181

<211> 14

<212> PRT

<213> Artificial Sequence

<220>

<223> Heavy Chain amino acid sequence

<400> 181

Glu Leu Gly Ser Gly Ser Tyr Tyr Pro Gly Tyr Phe Gln His
1 5 10

<210> 182

<211> 116

<212> PRT

<213> Artificial Sequence

<220>

<223> Light Chain amino acid sequence

<400> 182

Gln Asp Ile Gln Met Thr Gln Ser Pro Ala Thr Leu Tyr Val Ser Pro
1 5 10 15
Gly Glu Arg Ala Thr Leu Ser Cys Arg Ala Ser Gln Ser Val Ser Arg
20 25 30
Asn Leu Ala Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro Arg Leu Leu
35 40 45
Ile Tyr Gly Ala Ser Thr Arg Ala Thr Gly Ile Pro Ala Arg Phe Ser
50 55 60
Gly Ser Gly Ser Arg Thr Glu Phe Thr Leu Thr Ile Ser Ser Leu Gln
65 70 75 80
Ser Glu Asp Phe Ala Val Tyr His Cys Gln Gln Tyr Asn Ser Arg Pro
85 90 95
Leu Thr Phe Gly Gly Gly Thr Lys Val Glu Ile Lys Arg Thr Val Ala
100 105 110
Ala Pro Ser Val
115

<210> 183

<211> 123

<212> PRT

<213> Artificial Sequence

<220>

<223> Heavy Chain amino acid sequence

<400> 183

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1 5 10 15
Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Trp Tyr
20 25 30

Phe	Met	Asn	Trp	Val	Arg	Gln	Ala	Pro	Gly	Lys	Gly	Leu	Glu	Trp	Val
		35					40					45			
Ser	Ser	Ile	Tyr	Pro	Ser	Gly	Gly	Tyr	Thr	Met	Tyr	Ala	Asp	Ser	Val
	50					55					60				
Lys	Gly	Arg	Phe	Thr	Ile	Ser	Arg	Asp	Asn	Ser	Lys	Lys	Thr	Leu	Tyr
65					70				75					80	
Leu	Gln	Met	Asn	Ser	Leu	Arg	Ala	Glu	Asp	Thr	Ala	Val	Tyr	Tyr	Cys
			85						90					95	
Ala	Ser	Asp	Phe	Gly	Ser	Trp	Gly	Gln	Gly	Thr	Leu	Val	Thr	Val	Ser
		100						105					110		
Ser	Ala	Ser	Thr	Lys	Gly	Pro	Ser	Val	Phe	Pro					
		115					120								

<210> 184

<211> 11

<212> PRT

<213> Artificial Sequence

<220>

<223> Light Chain amino acid sequence

<400> 184

Arg	Ala	Ser	Gln	Ser	Val	Ser	Arg	Asn	Leu	Ala
1				5					10	

<210> 185

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> Light Chain amino acid sequence

<400> 185

Gly	Ala	Ser	Thr	Arg	Ala	Thr
1				5		

<210> 186

<211> 9

<212> PRT

<213> Artificial Sequence

<220>

<223> Light Chain amino acid sequence

<400> 186

Gln	Gln	Tyr	Asn	Ser	Arg	Pro	Leu	Thr
1				5				

<210> 187

<211> 5

<212> PRT

<213> Artificial Sequence

<220>

<223> Heavy Chain amino acid sequence

<400> 187

Trp Tyr Phe Met Asn
 1 5

<210> 188

<211> 17

<212> PRT

<213> Artificial Sequence

<220>

<223> Heavy Chain amino acid sequence

<400> 188

Ser Ile Tyr Pro Ser Gly Gly Tyr Thr Met Tyr Ala Asp Ser Val Lys
 1 5 10 15
 Gly

<210> 189

<211> 4

<212> PRT

<213> Artificial Sequence

<220>

<223> Heavy Chain amino acid sequence

<400> 189

Asp Phe Gly Ser
 1

<210> 190

<211> 119

<212> PRT

<213> Artificial Sequence

<220>

<223> Light Chain amino acid sequence

<400> 190

Gln Ser Ala Leu Thr Gln Pro Ala Ser Val Ser Gly Ser Pro Gly Gln
 1 5 10 15
 Ser Ile Thr Ile Ser Cys Thr Gly Thr Ser Ser Asp Val Gly Tyr Tyr
 20 25 30
 Asp Tyr Val Ser Trp Tyr Gln His His Pro Gly Lys Ala Pro Lys Leu
 35 40 45
 Ile Ile Tyr Asp Val Thr Ser Arg Pro Ser Gly Val Ser Ser His Phe
 50 55 60
 Ser Gly Ser Lys Ser Gly Asn Thr Ala Ser Leu Thr Ile Ser Gly Leu
 65 70 75 80
 Gln Ala Asp Asp Glu Ala Asp Tyr Tyr Cys Ser Ser Tyr Thr Ser Gly
 85 90 95
 Ser Thr Arg Tyr Val Phe Gly Pro Gly Thr Lys Val Thr Val Leu Gly
 100 105 110
 Gln Pro Lys Ala Asn Pro Thr
 115

<210> 191

<211> 131
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 191
 Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
 1 5 10 15
 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Asp Tyr
 20 25 30
 Tyr Met Arg Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
 35 40 45
 Ser Arg Ile Tyr Pro Ser Gly Gly His Thr Trp Tyr Ala Asp Ser Val
 50 55 60
 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
 65 70 75 80
 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
 85 90 95
 Ala Arg His Arg Ala Gly Ser Ser Gly Trp Tyr Ser Asp Tyr Trp Gly
 100 105 110
 Gln Gly Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser
 115 120 125
 Val Phe Pro
 130

<210> 192
 <211> 14
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 192
 Thr Gly Thr Ser Ser Asp Val Gly Tyr Tyr Asp Tyr Val Ser
 1 5 10

<210> 193
 <211> 7
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 193
 Asp Val Thr Ser Arg Pro Ser
 1 5

<210> 194
 <211> 11
 <212> PRT
 <213> Artificial Sequence

<220>

<223> Light Chain amino acid sequence

<400> 194

Ser Ser Tyr Thr Ser Gly Ser Thr Arg Tyr Val
1 5 10

<210> 195

<211> 5

<212> PRT

<213> Artificial Sequence

<220>

<223> Heavy Chain amino acid sequence

<400> 195

Asp Tyr Tyr Met Arg
1 5

<210> 196

<211> 17

<212> PRT

<213> Artificial Sequence

<220>

<223> Heavy Chain amino acid sequence

<400> 196

Arg Ile Tyr Pro Ser Gly Gly His Thr Trp Tyr Ala Asp Ser Val Lys
1 5 10 15
Gly

<210> 197

<211> 12

<212> PRT

<213> Artificial Sequence

<220>

<223> Heavy Chain amino acid sequence

<400> 197

His Arg Ala Gly Ser Ser Gly Trp Tyr Ser Asp Tyr
1 5 10

<210> 198

<211> 116

<212> PRT

<213> Artificial Sequence

<220>

<223> Light Chain amino acid sequence

<400> 198

Gln Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val
1 5 10 15
Gly Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Ser Ser
20 25 30


```

Tyr Leu Asn Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu
   35           40           45
Ile Tyr Ala Ala Ser Ser Leu Gln Ser Gly Val Pro Ser Arg Phe Ser
   50           55           60
Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln
  65           70           75           80
Pro Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Tyr Ser Thr Arg
           85           90           95
Trp Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys Arg Thr Val Ala
   100           105           110
Ala Pro Ser Val
      115

```

<210> 199

<211> 137

<212> PRT

<213> Artificial Sequence

<220>

<223> Heavy Chain amino acid sequence

<400> 199

```

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
  1           5           10           15
Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Thr Tyr
           20           25           30
Phe Met Arg Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
           35           40           45
Ser Tyr Ile Val Pro Ser Gly Gly Asn Thr Leu Tyr Ala Asp Ser Val
           50           55           60
Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
  65           70           75           80
Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
           85           90           95
Ala Arg Glu Glu Trp Asp Val Leu Leu Trp Phe Gly Glu Leu Ser Ala
           100           105           110
Ala Phe Asp Ile Trp Gly Gln Gly Thr Met Val Thr Val Ser Ser Ala
           115           120           125
Ser Thr Lys Gly Pro Ser Val Phe Pro
   130           135

```

<210> 200

<211> 11

<212> PRT

<213> Artificial Sequence

<220>

<223> Light Chain amino acid sequence

<400> 200

```

Arg Ala Ser Gln Ser Ile Ser Ser Tyr Leu Asn
  1           5           10

```

<210> 201

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> Light Chain amino acid sequence

<400> 201

Ala Ala Ser Ser Leu Gln Ser

1

5

<210> 202

<211> 9

<212> PRT

<213> Artificial Sequence

<220>

<223> Light Chain amino acid sequence

<400> 202

Gln Gln Ser Tyr Ser Thr Arg Trp Thr

1

5

<210> 203

<211> 5

<212> PRT

<213> Artificial Sequence

<220>

<223> Heavy Chain amino acid sequence

<400> 203

Thr Tyr Phe Met Arg

1

5

<210> 204

<211> 17

<212> PRT

<213> Artificial Sequence

<220>

<223> Heavy Chain amino acid sequence

<400> 204

Tyr Ile Val Pro Ser Gly Gly Asn Thr Leu Tyr Ala Asp Ser Val Lys

1

5

10

15

Gly

<210> 205

<211> 18

<212> PRT

<213> Artificial Sequence

<220>

<223> Heavy Chain amino acid sequence

<400> 205

Glu Glu Trp Asp Val Leu Leu Trp Phe Gly Glu Leu Ser Ala Ala Phe

1

5

10

15

Asp Ile

<210> 206

<211> 116

<212> PRT

<213> Artificial Sequence

<220>

<223> Light Chain amino acid sequence

<400> 206

[illegible]

<210> 207

<211> 132

<212> PRT

<213> Artificial Sequence

<220>

<223> Heavy Chain amino acid sequence

<400> 207

[illegible]

<210> 208
 <211> 11
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 208
 Arg Ala Ser Gln Gly Ile Arg His Tyr Leu Gly
 1 5 10

<210> 209
 <211> 7
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 209
 Ala Ala Ser Ser Leu Gln Phe
 1 5

<210> 210
 <211> 9
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 210
 Leu Gln His Asn Ser Phe Pro Pro Ala
 1 5

<210> 211
 <211> 5
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 211
 Pro Tyr Asp Met Trp
 1 5

<210> 212
 <211> 17
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 212

Tyr Ile Ser Ser Ser Gly Gly Lys Thr Met Tyr Ala Asp Ser Val Lys
 1 5 10 15
 Gly

<210> 213
 <211> 13
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 213
 Leu Gly Gly Asn Ser His Tyr Tyr Tyr Gly Met Asp Val
 1 5 10

<210> 214
 <211> 118
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 214
 Gln Ser Glu Leu Thr Gln Pro Pro Ser Ala Ser Ala Thr Pro Gly Gln
 1 5 10 15
 Arg Val Thr Ile Ser Cys Ser Gly Ser Ser Ser Asn Ile Gly Arg Asn
 20 25 30
 Leu Val Tyr Trp Tyr Gln Gln Leu Pro Gly Thr Ala Pro Lys Leu Leu
 35 40 45
 Ile Tyr Ser Asn Asn Gln Arg Pro Ser Gly Val Pro Asp Arg Phe Ser
 50 55 60
 Gly Ser Lys Ser Gly Thr Ser Ala Ser Leu Ala Ile Ser Gly Leu Arg
 65 70 75 80
 Ser Glu Glu Glu Ala Asp Tyr Tyr Cys Ala Ala Trp Asp Asp Ser Leu
 85 90 95
 Ser Gly Trp Val Phe Gly Gly Gly Thr Arg Leu Thr Val Leu Gly Gln
 100 105 110
 Pro Lys Ala Ala Pro Ser
 115

<210> 215
 <211> 131
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 215
 Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
 1 5 10 15
 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Trp Tyr
 20 25 30
 His Met Arg Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val

	35					40					45						
Ser	Ile	Tyr	Pro	Ser	Gly	Gly	Val	Thr	Ser	Tyr	Ala	Asp	Ser	Val	Lys		
	50					55					60						
Gly	Arg	Phe	Thr	Ile	Ser	Arg	Asp	Asn	Ser	Lys	Asn	Thr	Leu	Tyr	Leu		
65					70					75					80		
Gln	Met	Asn	Ser	Leu	Arg	Ala	Glu	Asp	Thr	Ala	Val	Tyr	Tyr	Cys	Ala		
				85					90					95			
Arg	Glu	Thr	Ser	Gly	Trp	Tyr	Arg	Asp	Arg	Trp	Phe	Asp	Pro	Trp	Gly		
			100					105					110				
Gln	Gly	Thr	Leu	Val	Thr	Val	Ser	Ala	Ser	Thr	Lys	Gly	Pro	Ser			
		115					120				125						
Val	Phe	Pro															
	130																

<210> 216

<211> 13

<212> PRT

<213> Artificial Sequence

<220>

<223> Light Chain amino acid sequence

<400> 216

Ser	Gly	Ser	Ser	Ser	Asn	Ile	Gly	Arg	Asn	Leu	Val	Tyr
1				5					10			

<210> 217

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> Light Chain amino acid sequence

<400> 217

Ser	Asn	Asn	Gln	Arg	Pro	Ser
1			5			

<210> 218

<211> 11

<212> PRT

<213> Artificial Sequence

<220>

<223> Light Chain amino acid sequence

<400> 218

Ala	Ala	Trp	Asp	Asp	Ser	Leu	Ser	Gly	Trp	Val
1				5					10	

<210> 219

<211> 5

<212> PRT

<213> Artificial Sequence

<220>

<223> Heavy Chain amino acid sequence

<400> 219

Trp Tyr His Met Arg
1 5

<210> 220

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Heavy Chain amino acid sequence

<400> 220

Ile Tyr Pro Ser Gly Gly Val Thr Asp Tyr Ala Asp Ser Val Lys Gly
1 5 10 15

<210> 221

<211> 13

<212> PRT

<213> Artificial Sequence

<220>

<223> Heavy Chain amino acid sequence

<400> 221

Glu Thr Ser Gly Trp Tyr Arg Asp Arg Trp Phe Asp Pro
1 5 10

<210> 222

<211> 119

<212> PRT

<213> Artificial Sequence

<220>

<223> Light Chain amino acid sequence

<400> 222

Gln Ser Val Leu Thr Gln Thr Ala Ser Val Ser Gly Ser Pro Gly Gln
1 5 10 15
Ser Ile Thr Ile Ser Cys Thr Gly Thr Ser Ser Asp Ile Gly Asp Tyr
20 25 30
Glu Tyr Val Ser Trp Tyr Gln Gln His Pro Gly Lys Ala Pro Lys Val
35 40 45
Ile Leu Tyr Glu Val Ser Asn Arg Pro Ser Gly Val Pro Asp Arg Phe
50 55 60
Ser Gly Ser Lys Ser Gly Asn Thr Ala Ser Leu Thr Ile Ser Gly Leu
65 70 75 80
Gln Ala Glu Asp Glu Ala Asp Tyr Tyr Cys Gly Ser Tyr Arg Lys Ser
85 90 95
Ser Thr Pro Tyr Val Phe Gly Thr Gly Thr Lys Val Ser Val Leu Gly
100 105 110
Gln Pro Lys Ala Asn Pro Thr
115

<210> 223

<211> 138

<212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 223
 Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
 1 5 10 15
 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Tyr Tyr
 20 25 30
 His Met Trp Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
 35 40 45
 Ser Val Ile Val Pro Ser Gly Gly Gly Thr Gln Tyr Ala Asp Ser Val
 50 55 60
 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
 65 70 75 80
 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
 85 90 95
 Ala Arg Asp Gly His Ser Ser Ser Trp Tyr Gly Gly Gly Ala His Tyr
 100 105 110
 Tyr Gly Met Asp Val Trp Gly Gln Gly Thr Thr Val Thr Val Ser Ser
 115 120 125
 Ala Ser Thr Lys Gly Pro Ser Val Phe Pro
 130 135

<210> 224
 <211> 14
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 224
 Thr Gly Thr Ser Ser Asp Ile Gly Asp Tyr Glu Tyr Val Ser
 1 5 10

<210> 225
 <211> 8
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 225
 Tyr Glu Val Ser Asn Arg Pro Ser
 1 5

<210> 226
 <211> 11
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 226
 Gly Ser Tyr Arg Lys Ser Ser Thr Pro Tyr Val
 1 5 10

<210> 227
 <211> 5
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 227
 Tyr Tyr His Met Trp
 1 5

<210> 228
 <211> 17
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 228
 Val Ile Val Pro Ser Gly Gly Gly Thr Gln Tyr Ala Asp Ser Val Lys
 1 5 10 15
 Gly

<210> 229
 <211> 19
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 229
 Asp Gly His Ser Ser Ser Trp Tyr Gly Gly Gly Ala His Tyr Tyr Gly
 1 5 10 15
 Met Asp Val

<210> 230
 <211> 116
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 230
 Gln Asp Ile Gln Met Thr Gln Ser Pro Ala Thr Leu Ser Leu Ser Pro
 1 5 10 15
 Gly Glu Arg Ala Thr Leu Ser Cys Arg Ala Ser Gln Ser Val Ser Ser

		20						25					30				
Tyr	Leu	Ala	Trp	Tyr	Gln	Gln	Lys	Pro	Gly	Gln	Ala	Pro	Arg	Leu	Leu		
		35					40					45					
Ile	Tyr	Gly	Ala	Ser	Ser	Arg	Ala	Thr	Gly	Ile	Pro	Asp	Arg	Phe	Ser		
	50					55					60						
Gly	Ser	Gly	Ser	Gly	Thr	Asp	Phe	Thr	Leu	Thr	Ile	Gly	Arg	Leu	Glu		
65					70					75					80		
Pro	Glu	Asp	Phe	Ala	Val	Tyr	Tyr	Cys	Gln	Gln	Tyr	Ser	Ser	Ser	Pro		
				85					90					95			
Val	Thr	Phe	Gly	Gln	Gly	Thr	Arg	Leu	Glu	Ile	Lys	Arg	Thr	Val	Ala		
			100					105					110				
Ala	Pro	Ser	Val														
		115															

<210> 231

<211> 123

<212> PRT

<213> Artificial Sequence

<220>

<223> Heavy Chain amino acid sequence

<400> 231

Glu	Val	Gln	Leu	Leu	Glu	Ser	Gly	Gly	Gly	Leu	Val	Gln	Pro	Gly	Gly		
1				5					10					15			
Ser	Leu	Arg	Leu	Ser	Cys	Ala	Ala	Ser	Gly	Phe	Thr	Phe	Ser	Ser	Tyr		
			20					25					30				
Arg	Met	Asn	Trp	Val	Arg	Gln	Ala	Pro	Gly	Lys	Gly	Leu	Glu	Trp	Val		
		35				40						45					
Ser	Gly	Ile	Val	Pro	Ser	Gly	Gly	Lys	Thr	Phe	Tyr	Ala	Asp	Ser	Val		
	50					55					60						
Lys	Gly	Arg	Phe	Thr	Ile	Ser	Arg	Asp	Asn	Ser	Lys	Asn	Thr	Leu	Tyr		
65					70					75					80		
Leu	Gln	Met	Asn	Ser	Leu	Arg	Ala	Glu	Asp	Thr	Ala	Val	Tyr	Tyr	Cys		
			85					90						95			
Ala	Ser	Asp	Phe	Gly	Ser	Trp	Gly	Gln	Gly	Thr	Leu	Val	Thr	Val	Ser		
			100					105					110				
Ser	Ala	Ser	Thr	Lys	Gly	Pro	Ser	Val	Phe	Pro							
		115					120										

<210> 232

<211> 11

<212> PRT

<213> Artificial Sequence

<220>

<223> Light Chain amino acid sequence

<400> 232

Arg	Ala	Ser	Gln	Ser	Val	Ser	Ser	Tyr	Leu	Ala							
1				5					10								

<210> 233

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> Light Chain amino acid sequence

<400> 233

Gly Ala Ser Ser Arg Ala Thr

1

5

<210> 234

<211> 9

<212> PRT

<213> Artificial Sequence

<220>

<223> Light Chain amino acid sequence

<400> 234

Gln Gln Tyr Ser Ser Ser Pro Val Thr

1

5

<210> 235

<211> 5

<212> PRT

<213> Artificial Sequence

<220>

<223> Heavy Chain amino acid sequence

<400> 235

Ser Tyr Arg Met Asn

1

5

<210> 236

<211> 17

<212> PRT

<213> Artificial Sequence

<220>

<223> Heavy Chain amino acid sequence

<400> 236

Gly Ile Val Pro Ser Gly Gly Lys Thr Phe Tyr Ala Asp Ser Val Lys

1

5

10

15

Gly

<210> 237

<211> 4

<212> PRT

<213> Artificial Sequence

<220>

<223> Heavy Chain amino acid sequence

<400> 237

Asp Phe Gly Ser

1

<210> 238
 <211> 116
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 238
 Gln Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val
 1 5 10 15
 Gly Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Arg Ile Ser Ser
 20 25 30
 Tyr Val Asn Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu
 35 40 45
 Ile Tyr Ser Ala Ser Ser Leu Gln Ser Gly Val Pro Ser Arg Phe Ser
 50 55 60
 Gly Ser Val Ser Gly Thr Glu Phe Thr Leu Thr Ile Ser Ser Leu Gln
 65 70 75 80
 Pro Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Tyr Arg Thr Pro
 85 90 95
 Pro Phe Phe Gly Gln Gly Thr Lys Leu Glu Val Lys Arg Thr Val Ala
 100 105 110
 Ala Pro Ser Val
 115

<210> 239
 <211> 129
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 239
 Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
 1 5 10 15
 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Leu Tyr
 20 25 30
 Gln Met Leu Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
 35 40 45
 Ser Gly Ile Val Ser Ser Gly Gly Leu Thr Gly Tyr Ala Asp Ser Val
 50 55 60
 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
 65 70 75 80
 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
 85 90 95
 Ala Arg His Asn Arg Ala Ile Gly Thr Phe Asp Tyr Trp Gly Gln Gly
 100 105 110
 Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val Phe
 115 120 125
 Pro

<210> 240
 <211> 11
 <212> PRT

<213> Artificial Sequence

<220>

<223> Light Chain amino acid sequence

<400> 240

Arg Ala Ser Gln Arg Ile Ser Ser Tyr Val Asn
1 5 10

<210> 241

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> Light Chain amino acid sequence

<400> 241

Ser Ala Ser Ser Leu Gln Ser
1 5

<210> 242

<211> 9

<212> PRT

<213> Artificial Sequence

<220>

<223> Light Chain amino acid sequence

<400> 242

Gln Gln Ser Tyr Arg Thr Pro Pro Phe
1 5

<210> 243

<211> 5

<212> PRT

<213> Artificial Sequence

<220>

<223> Heavy Chain amino acid sequence

<400> 243

Leu Tyr Gln Met Leu
1 5

<210> 244

<211> 17

<212> PRT

<213> Artificial Sequence

<220>

<223> Heavy Chain amino acid sequence

<400> 244

Gly Ile Val Ser Ser Gly Gly Leu Thr Gly Tyr Ala Asp Ser Val Lys
1 5 10 15
Gly

<210> 245
 <211> 10
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 245
 His Asn Arg Ala Ile Gly Thr Phe Asp Tyr
 1 5 10

<210> 246
 <211> 115
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 246
 Gln Asp Ile Gln Met Thr Gln Ser Pro Ala Thr Leu Ser Leu Ser Pro
 1 5 10 15
 Gly Glu Arg Ala Thr Leu Ser Cys Arg Ala Ser Gln Ser Val Ser Arg
 20 25 30
 Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro Arg Leu Leu
 35 40 45
 Ile Tyr Gly Ala Ser Thr Arg Ala Thr Gly Ile Pro Ala Arg Phe Ser
 50 55 60
 Gly Ser Gly Ser Gly Thr Glu Phe Thr Leu Thr Ile Ser Ser Leu Gln
 65 70 75 80
 Ser Glu Asp Phe Ala Val Tyr Tyr Cys Gln Gln Tyr Asn Asn Trp Pro
 85 90 95
 Ser Phe Gly Gly Thr Lys Val Glu Ile Lys Arg Thr Val Ala Ala
 100 105 110
 Pro Ser Val
 115

<210> 247
 <211> 123
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 247
 Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
 1 5 10 15
 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Asn Tyr
 20 25 30
 Ser Met Asp Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
 35 40 45
 Ser Trp Ile Ser Pro Ser Gly Gly Leu Thr Thr Tyr Ala Asp Ser Val
 50 55 60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
 65 70 75 80
 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
 85 90 95
 Ala Ser Asp Phe Gly Ser Trp Gly Gln Gly Thr Leu Val Thr Val Ser
 100 105 110
 Ser Ala Ser Thr Lys Gly Pro Ser Val Phe Pro
 115 120

<210> 248
 <211> 11
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 248
 Arg Ala Ser Gln Ser Val Ser Arg Tyr Leu Ala
 1 5 10

<210> 249
 <211> 7
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 249
 Gly Ala Ser Thr Arg Ala Thr
 1 5

<210> 250
 <211> 8
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 250
 Gln Gln Tyr Asn Asn Trp Pro Ser
 1 5

<210> 251
 <211> 5
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 251
 Asn Tyr Ser Met Asp
 1 5

<210> 252
 <211> 17
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 252
 Trp Ile Ser Pro Ser Gly Gly Leu Thr Thr Tyr Ala Asp Ser Val Lys
 1 5 10 15
 Gly

<210> 253
 <211> 4
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 253
 Asp Phe Gly Ser
 1

<210> 254
 <211> 124
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 254
 Gln Ser Val Leu Thr Gln Pro Pro Tyr Ala Ser Ala Ser Leu Gly Ala
 1 5 10 15
 Ser Val Thr Leu Thr Cys Thr Leu Ser Ser Gly Tyr Ser Asn Tyr Lys
 20 25 30
 Val Asp Trp Tyr Gln Gln Arg Pro Gly Lys Gly Pro Gln Phe Val Met
 35 40 45
 Arg Val Gly Ser Gly Gly Ile Val Gly Ser Lys Gly Asp Gly Ile Pro
 50 55 60
 Asp Arg Phe Ser Val Leu Gly Ser Gly Leu Tyr Arg Tyr Leu Thr Ile
 65 70 75 80
 Lys Asn Ile Gln Glu Glu Asp Glu Ser Asp Tyr Tyr Cys Gly Ala Asp
 85 90 95
 His Gly Arg Gly Gly Thr Phe Val Trp Val Phe Gly Gly Gly Thr Lys
 100 105 110
 Leu Thr Val Leu Gly Gln Pro Lys Ala Ala Pro Ser
 115 120

<210> 255
 <211> 136
 <212> PRT
 <213> Artificial Sequence

<220>

<223> Heavy Chain amino acid sequence

<400> 255

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Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
 1             5             10             15
Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Tyr Lys
      20             25             30
Met Met Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val Ser
      35             40             45
Tyr Ile Ser Ser Ser Gly Gly Ile Thr Thr Tyr Ala Asp Ser Val Lys
      50             55             60
Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr Leu
65             70             75             80
Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys Ala
      85             90             95
Arg Asp Pro Thr Tyr Asp Phe Trp Ser Gly Tyr Tyr Tyr Tyr Tyr Tyr
      100            105            110
Met Asp Val Trp Gly Lys Gly Thr Thr Val Thr Val Ser Ser Ala Ser
      115            120            125
Thr Lys Gly Pro Ser Val Phe Pro
      130            135

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<210> 256

<211> 12

<212> PRT

<213> Artificial Sequence

<220>

<223> Light Chain amino acid sequence

<400> 256

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Thr Leu Ser Ser Gly Tyr Ser Asn Tyr Lys Val Asp
 1             5             10

```

<210> 257

<211> 13

<212> PRT

<213> Artificial Sequence

<220>

<223> Light Chain amino acid sequence

<400> 257

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Arg Val Gly Ser Gly Gly Ile Val Gly Ser Lys Gly Asp
 1             5             10

```

<210> 258

<211> 13

<212> PRT

<213> Artificial Sequence

<220>

<223> Light Chain amino acid sequence

<400> 258

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Gly Ala Asp His Gly Arg Gly Gly Thr Phe Val Trp Val

```

1 5 10

<210> 259
 <211> 5
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 259
 Ser Tyr Lys Met Met
 1 5

<210> 260
 <211> 17
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 260
 Tyr Ile Ser Ser Ser Gly Gly Ile Thr Thr Tyr Ala Asp Ser Val Lys
 1 5 10 15
 Gly

<210> 261
 <211> 19
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 261
 Arg Asp Pro Thr Tyr Asp Phe Trp Ser Gly Tyr Tyr Tyr Tyr Tyr Tyr
 1 5 10 15
 Met Asp Val

<210> 262
 <211> 118
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 262
 Gln Ser Ala Leu Thr Gln Pro Ser Ser Ala Ser Gly Thr Pro Gly Gln
 1 5 10 15
 Arg Val Ser Ile Ser Cys Ser Gly Ser Ser Tyr Asn Ile Gly Val Tyr
 20 25 30
 Asp Val Tyr Trp Tyr Gln Gln Leu Pro Gly Thr Ala Pro Lys Leu Leu
 35 40 45

Ile Tyr Thr Asn Asn Gln Arg Pro Ser Gly Val Pro Asp Arg Phe Ser
 50 55 60
 Gly Ser Lys Ser Gly Thr Ser Ala Ser Leu Ala Ile Ser Gly Leu Gln
 65 70 75 80
 Ser Glu Asp Glu Ala Asp Tyr Tyr Cys Ala Ala Trp Asp Asp Ser Leu
 85 90 95
 Ser Gly Trp Val Phe Gly Gly Gly Thr Lys Val Thr Val Leu Gly Gln
 100 105 110
 Pro Lys Ala Ala Pro Ser
 115

<210> 263
 <211> 137
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 263
 Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
 1 5 10 15
 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Gln Tyr
 20 25 30
 Asn Met Pro Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
 35 40 45
 Ser Ser Ile Val Pro Ser Gly Gly Phe Thr Ala Tyr Ala Asp Ser Val
 50 55 60
 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
 65 70 75 80
 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
 85 90 95
 Ala Arg Val Asp Cys Ser Gly Gly Ser Cys Tyr Arg Gly Pro Gln Asn
 100 105 110
 Tyr Phe Asp Tyr Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser Ala
 115 120 125
 Ser Thr Lys Gly Pro Ser Val Phe Pro
 130 135

<210> 264
 <211> 13
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 264
 Ser Gly Ser Ser Tyr Asn Ile Gly Val Tyr Asp Val Tyr
 1 5 10

<210> 265
 <211> 7
 <212> PRT
 <213> Artificial Sequence

<220>

<223> Light Chain amino acid sequence

<400> 265

Thr Asn Asn Gln Arg Pro Ser
1 5

<210> 266

<211> 11

<212> PRT

<213> Artificial Sequence

<220>

<223> Light Chain amino acid sequence

<400> 266

Ala Ala Trp Asp Asp Ser Leu Ser Gly Trp Val
1 5 10

<210> 267

<211> 5

<212> PRT

<213> Artificial Sequence

<220>

<223> Light Chain amino acid sequence

<400> 267

Gln Tyr Asn Met Pro
1 5

<210> 268

<211> 17

<212> PRT

<213> Artificial Sequence

<220>

<223> Heavy Chain amino acid sequence

<400> 268

Ser Ile Val Pro Ser Gly Gly Phe Thr Ala Tyr Ala Asp Ser Val Lys
1 5 10 15
Gly

<210> 269

<211> 18

<212> PRT

<213> Artificial Sequence

<220>

<223> Light Chain amino acid sequence

<400> 269

Val Asp Cys Ser Gly Gly Ser Cys Tyr Arg Gly Pro Gln Asn Tyr Phe
1 5 10 15
Asp Tyr

<210> 270
 <211> 119
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 270
 Gln Tyr Glu Leu Thr Gln Pro Ala Ser Val Ser Gly Ser Pro Gly Gln
 1 5 10 15
 Ser Ile Thr Ile Ser Cys Thr Gly Thr Ser Ser Asp Val Gly Gly Tyr
 20 25 30
 Asn Tyr Val Ser Trp Tyr Gln Gln His Pro Gly Lys Ala Pro Lys Leu
 35 40 45
 Met Ile Tyr Glu Val Ser Asn Arg Pro Ser Gly Val Ser Asn Arg Phe
 50 55 60
 Ser Gly Ser Lys Ser Asp Asn Thr Ala Ser Leu Thr Ile Ser Gly Leu
 65 70 75 80
 Gln Ala Glu Asp Glu Ala Asp Tyr Tyr Cys Gly Ser Tyr Arg Lys Ser
 85 90 95
 Ser Thr Pro Tyr Val Phe Gly Thr Gly Thr Lys Val Ser Val Leu Gly
 100 105 110
 Gln Pro Lys Ala Asn Pro Thr
 115

<210> 271
 <211> 135
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 271
 Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
 1 5 10 15
 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Gln Tyr
 20 25 30
 Met Met Thr Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
 35 40 45
 Ser Tyr Ile Gly Ser Ser Gly Gly Gln Thr Lys Tyr Ala Asp Ser Val
 50 55 60
 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
 65 70 75 80
 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
 85 90 95
 Ala Arg Asp Pro Gly Val Ala Val Ala Gly Tyr Tyr Tyr Tyr Gly Met
 100 105 110
 Asp Val Trp Gly Gln Gly Thr Thr Val Thr Val Ser Ser Ala Ser Thr
 115 120 125
 Lys Gly Pro Ser Val Phe Pro
 130 135

<210> 272
 <211> 14

<212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 272
 Thr Gly Thr Ser Ser Asp Val Gly Gly Tyr Asn Tyr Val Ser
 1 5 10

<210> 273
 <211> 7
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 273
 Glu Val Ser Asn Arg Pro Ser
 1 5

<210> 274
 <211> 11
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 274
 Gly Ser Tyr Arg Lys Ser Ser Thr Pro Tyr Val
 1 5 10

<210> 275
 <211> 5
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 275
 Gln Tyr Met Met Thr
 1 5

<210> 276
 <211> 17
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 276
 Tyr Ile Gly Ser Ser Gly Gly Gln Thr Lys Tyr Ala Asp Ser Val Lys
 1 5 10 15

Gly

<210> 277
 <211> 16
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 277
 Asp Pro Gly Val Ala Val Ala Gly Tyr Tyr Tyr Tyr Gly Met Asp Val
 1 5 10 15

<210> 278
 <211> 116
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 278
 Gln Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Val Ser Ala Ser Val
 1 5 10 15
 Gly Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Arg Gly Ile Ser Arg
 20 25 30
 Trp Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu
 35 40 45
 Ile Tyr Gly Ala Ser Thr Leu Gln Lys Gly Val Pro Ser Arg Phe Thr
 50 55 60
 Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Thr Ser Leu Gln
 65 70 75 80
 Pro Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Gly Asn Ser Phe Pro
 85 90 95
 Phe Thr Phe Gly Pro Gly Thr Lys Val Asp Ile Lys Arg Thr Val Ala
 100 105 110
 Ala Pro Ser Val
 115

<210> 279
 <211> 132
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 279
 Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
 1 5 10 15
 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Gly Tyr
 20 25 30
 Trp Met Ser Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
 35 40 45
 Ser Val Ile Arg Pro Ser Gly Gly Lys Thr Gly Tyr Ala Asp Ser Val

50		55		60	
Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Phe Lys Asn Thr Leu Tyr					
65		70		75	80
Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys					
	85		90		95
Ala Arg Val Arg Ala Pro Gly Tyr Tyr Tyr Tyr Gly Met Asp Val Trp					
	100		105		110
Gly Gln Gly Thr Thr Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro					
	115		120		125
Ser Val Phe Pro					
130					

<210> 280
 <211> 11
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 280
 Arg Ala Ser Arg Gly Ile Ser Arg Trp Leu Ala
 1 5 10

<210> 281
 <211> 7
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 281
 Gly Ala Ser Thr Leu Gln Lys
 1 5

<210> 282
 <211> 9
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 282
 Gln Gln Gly Asn Ser Phe Pro Phe Thr
 1 5

<210> 283
 <211> 5
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 283

Gly Tyr Trp Met Ser
1 5

<210> 284
<211> 17
<212> PRT
<213> Artificial Sequence

<220>
<223> Heavy Chain amino acid sequence

<400> 284
Val Ile Arg Pro Ser Gly Gly Lys Thr Gly Tyr Ala Asp Ser Val Lys
1 5 10 15
Gly

<210> 285
<211> 13
<212> PRT
<213> Artificial Sequence

<220>
<223> Heavy Chain amino acid sequence

<400> 285
Val Arg Ala Pro Gly Tyr Tyr Tyr Tyr Gly Met Asp Val
1 5 10

<210> 286
<211> 119
<212> PRT
<213> Artificial Sequence

<220>
<223> Light Chain amino acid sequence

<400> 286
Gln Ser Val Leu Thr Gln Thr Ala Ser Val Ser Gly Ser Pro Gly Gln
1 5 10 15
Ser Ile Thr Ile Ser Cys Thr Gly Thr Ser Ser Asp Ile Gly Asp Tyr
20 25 30
Glu Tyr Val Ser Trp Tyr Gln Gln His Pro Gly Lys Ala Pro Lys Val
35 40 45
Ile Leu Tyr Glu Val Ser Asn Arg Pro Ser Gly Val Pro Asp Arg Phe
50 55 60
Ser Gly Ser Lys Ser Gly Asn Thr Ala Ser Leu Thr Ile Ser Gly Leu
65 70 75 80
Gln Ala Glu Asp Glu Ala Asp Tyr Tyr Cys Gly Ser Tyr Arg Lys Ser
85 90 95
Ser Thr Pro Tyr Val Phe Gly Thr Gly Thr Lys Val Ser Val Leu Gly
100 105 110
Gln Pro Lys Ala Asn Pro Thr
115

<210> 287
<211> 138

<212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 287
 Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
 1 5 10 15
 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Tyr Tyr
 20 25 30
 His Met Trp Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
 35 40 45
 Ser Val Ile Val Pro Ser Gly Gly Gly Thr Gln Tyr Ala Asp Ser Val
 50 55 60
 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
 65 70 75 80
 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
 85 90 95
 Ala Arg Asp Gly His Ser Ser Ser Trp Tyr Gly Gly Gly Ala His Tyr
 100 105 110
 Tyr Gly Met Asp Val Trp Gly Gln Gly Thr Thr Val Thr Val Ser Ser
 115 120 125
 Ala Ser Thr Lys Gly Pro Ser Val Phe Pro
 130 135

<210> 288
 <211> 14
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 288
 Thr Gly Thr Ser Ser Asp Ile Gly Asp Tyr Glu Tyr Val Ser
 1 5 10

<210> 289
 <211> 8
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 289
 Tyr Glu Val Ser Asn Arg Pro Ser
 1 5

<210> 290
 <211> 11
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 290
 Gly Ser Tyr Arg Lys Ser Ser Thr Pro Tyr Val
 1 5 10

<210> 291
 <211> 5
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 291
 Tyr Tyr His Met Trp
 1 5

<210> 292
 <211> 17
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 292
 Val Ile Val Pro Ser Gly Gly Gly Thr Gln Tyr Ala Asp Ser Val Lys
 1 5 10 15
 Gly

<210> 293
 <211> 19
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heavy Chain amino acid sequence

<400> 293
 Asp Gly His Ser Ser Ser Trp Tyr Gly Gly Gly Ala His Tyr Tyr Gly
 1 5 10 15
 Met Asp Val

<210> 294
 <211> 116
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Light Chain amino acid sequence

<400> 294
 Gln Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val
 1 5 10 15
 Gly Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Gly Ile Arg Asn

[illegible]

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<210> 295
<211> 132
<212> PRT
<213> Artificial Sequence
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<220>
<223> Heavy Chain amino acid sequence

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<400> 295
Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
 1          5          10          15
Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Phe Tyr
          20          25          30
Gly Met Pro Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
          35          40          45
Ser Gly Ile Tyr Pro Ser Gly Gly Val Thr Arg Tyr Ala Asp Ser Val
          50          55          60
Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65          70          75          80
Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
          85          90          95
Ala Lys Thr Tyr Ser Ser Ser Trp Tyr Gly Trp Tyr Phe Asp Tyr Trp
          100          105          110
Gly Gln Gly Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro
          115          120          125
Ser Val Phe Pro
          130

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<210> 296
<211> 11
<212> PRT
<213> Artificial Sequence
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<220>
<223> Light Chain amino acid sequence

<400> 296
Arg Ala Ser Gln Gly Ile Arg Asn Asp Leu Gly
1 5 10

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<210> 297
<211> 7
<212> PRT
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<213> Artificial Sequence

<220>

<223> Light Chain amino acid sequence

<400> 297

Gly Ala Ser Thr Leu Gln Ser

1 5

<210> 298

<211> 9

<212> PRT

<213> Artificial Sequence

<220>

<223> Light Chain amino acid sequence

<400> 298

Leu Gln Asp Tyr Asn Tyr Pro Tyr Thr

1 5

<210> 299

<211> 5

<212> PRT

<213> Artificial Sequence

<220>

<223> Heavy Chain amino acid sequence

<400> 299

Phe Tyr Gly Met Pro

1 5

<210> 300

<211> 17

<212> PRT

<213> Artificial Sequence

<220>

<223> Heavy Chain amino acid sequence

<400> 300

Gly Ile Tyr Pro Ser Gly Gly Val Thr Arg Tyr Ala Asp Ser Val Lys

1 5 10 15

Gly

<210> 301

<211> 13

<212> PRT

<213> Unknown

<220>

<223> Heavy Chain amino acid sequence

<400> 301

Thr Tyr Ser Ser Ser Trp Tyr Gly Trp Tyr Phe Asp Tyr

1 5 10

<210> 302
 <211> 117
 <212> PRT
 <213> Unknown

<220>
 <223> Light Chain amino acid sequence

<400> 302
 Gln Asp Ile Gln Met Thr Gln Ser Pro Gly Thr Leu Ser Leu Ser Pro
 1 5 10 15
 Gly Glu Arg Ala Thr Leu Ser Cys Arg Ala Ser Gln Ser Val Ser Ser
 20 25 30
 Ser Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro Arg Leu
 35 40 45
 Leu Ile Tyr Gly Ala Ser Ser Arg Ala Thr Gly Ile Pro Asp Arg Phe
 50 55 60
 Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Arg Leu
 65 70 75 80
 Glu Pro Glu Asp Phe Ala Val Tyr Tyr Cys Gln Gln Tyr Gly Ser Ser
 85 90 95
 Pro Trp Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys Arg Thr Val
 100 105 110
 Ala Ala Pro Ser Val
 115

<210> 303
 <211> 130
 <212> PRT
 <213> Unknown

<220>
 <223> Heavy Chain amino acid sequence

<400> 303
 Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
 1 5 10 15
 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Phe Tyr
 20 25 30
 Pro Met Pro Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
 35 40 45
 Ser Tyr Ile Ser Pro Ser Gly Gly Asp Thr Thr Tyr Ala Asp Ser Val
 50 55 60
 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Phe Tyr
 65 70 75 80
 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
 85 90 95
 Ala Arg Gly Gly Ser Tyr Ser Ser Ser Trp Tyr Gly Tyr Trp Gly Gln
 100 105 110
 Gly Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val
 115 120 125
 Phe Pro
 130

<210> 304

<211> 12
 <212> PRT
 <213> Unknown

<220>
 <223> Light Chain amino acid sequence

<400> 304
 Arg Ala Ser Gln Ser Val Ser Ser Ser Tyr Leu Ala
 1 5 10

<210> 305
 <211> 7
 <212> PRT
 <213> Unknown

<220>
 <223> Light Chain amino acid sequence

<400> 305
 Gly Ala Ser Ser Arg Ala Thr
 1 5

<210> 306
 <211> 9
 <212> PRT
 <213> Unknown

<220>
 <223> Light Chain amino acid sequence

<400> 306
 Gln Gln Tyr Gly Ser Ser Pro Trp Thr
 1 5

<210> 307
 <211> 5
 <212> PRT
 <213> Unknown

<220>
 <223> Heavy Chain amino acid sequence

<400> 307
 Phe Tyr Pro Met Pro
 1 5

<210> 308
 <211> 17
 <212> PRT
 <213> Unknown

<220>
 <223> Heavy Chain amino acid sequence

<400> 308
 Tyr Ile Ser Pro Ser Gly Gly Asp Thr Thr Tyr Ala Asp Ser Val Lys

1	5	10	15
Gly			

```
<210> 309
<211> 11
<212> PRT
<213> Unknown
```

<220>
<223> Heavy Chain amino acid sequence

<400> 309
Gly Gly Ser Tyr Ser Ser Ser Trp Tyr Gly Tyr
1 5 10

```
<210> 310
<211> 116
<212> PRT
<213> Unknown
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<220>
<223> Light Chain amino acid sequence

<400>	310
Gln Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Val Ser Ala Ser Val	
1 5 10 15	
Gly Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Arg Gly Ile Ser Arg	
20 25 30	
Trp Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu	
35 40 45	
Ile Tyr Gly Ala Ser Thr Leu Gln Lys Gly Val Pro Ser Arg Phe Thr	
50 55 60	
Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Thr Ser Leu Gln	
65 70 75 80	
Pro Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Gly Asn Ser Phe Pro	
85 90 95	
Phe Thr Phe Gly Pro Gly Thr Lys Val Asp Ile Lys Arg Thr Val Ala	
100 105 110	
Ala Pro Ser Val	
115	

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<210> 311
<211> 132
<212> PRT
<213> Unknown
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<220>
<223> Heavy Chain amino acid sequence

<400> 311
Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1 5 10 15
Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Gly Tyr
20 25 30
Trp Met Ser Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45

Ser	Val	Ile	Arg	Pro	Ser	Gly	Gly	Lys	Thr	Gly	Tyr	Ala	Asp	Ser	Val
50						55					60				
Lys	Gly	Arg	Phe	Thr	Ile	Ser	Arg	Asp	Asn	Phe	Lys	Asn	Thr	Leu	Tyr
65					70					75				80	
Leu	Gln	Met	Asn	Ser	Leu	Arg	Ala	Glu	Asp	Thr	Ala	Val	Tyr	Tyr	Cys
			85						90				95		
Ala	Arg	Val	Arg	Ala	Pro	Gly	Tyr	Tyr	Tyr	Tyr	Gly	Met	Asp	Val	Trp
			100					105					110		
Gly	Gln	Gly	Thr	Thr	Val	Thr	Val	Ser	Ser	Ala	Ser	Thr	Lys	Gly	Pro
		115					120						125		
Ser	Val	Phe	Pro												
130															

<210> 312
 <211> 11
 <212> PRT
 <213> Unknown

<220>
 <223> Light Chain amino acid sequence

<400> 312
Arg Ala Ser Arg Gly Ile Ser Arg Trp Leu Ala
1 5 10

<210> 313
 <211> 7
 <212> PRT
 <213> Unknown

<220>
 <223> Light Chain amino acid sequence

<400> 313
Gly Ala Ser Thr Leu Gln Lys
1 5

<210> 314
 <211> 9
 <212> PRT
 <213> Unknown

<220>
 <223> Light Chain amino acid sequence

<400> 314
Gln Gln Gly Asn Ser Phe Pro Phe Thr
1 5

<210> 315
 <211> 5
 <212> PRT
 <213> Unknown

<220>
 <223> Heavy Chain amino acid sequence

<400> 315

Gly Tyr Trp Met Ser
 1 5

<210> 316

<211> 17

<212> PRT

<213> Unknown

<220>

<223> Heavy Chain amino acid sequence

<400> 316

Val Ile Arg Pro Ser Gly Gly Lys Thr Gly Tyr Ala Asp Ser Val Lys
 1 5 10 15
 Gly

<210> 317

<211> 13

<212> PRT

<213> Unknown

<220>

<223> Heavy Chain amino acid sequence

<400> 317

Val Arg Ala Pro Gly Tyr Tyr Tyr Tyr Gly Met Asp Val
 1 5 10

<210> 318

<211> 118

<212> PRT

<213> Unknown

<220>

<223> Light Chain amino acid sequence

<400> 318

Gln Ser Val Leu Thr Gln Pro Ala Ser Val Ser Gly Ser Pro Gly Gln
 1 5 10 15
 Ser Ile Thr Ile Ser Cys Thr Gly Thr Ser Ser Asp Val Gly Gly Tyr
 20 25 30
 Asn Tyr Val Ser Trp Tyr Gln Arg His Pro Gly Lys Ala Pro Lys Leu
 35 40 45
 Ile Ile Tyr Asp Val Thr Asn Arg Pro Ser Gly Ala Ser Arg His Phe
 50 55 60
 Ser Gly Ser Lys Ser Gly Asn Thr Ala Ser Leu Thr Ile Ser Gly Leu
 65 70 75 80
 Gln Ala Asp Asp Glu Ala Asp Tyr Tyr Cys Val Ser Phe Thr Asn Ser
 85 90 95
 Asn Thr Phe Val Phe Gly Ser Gly Thr Arg Val Thr Val Leu Gly Gln
 100 105 110
 Pro Lys Ala Asn Pro Thr
 115

<210> 319

<211> 138
 <212> PRT
 <213> Unknown

<220>

<223> Heavy Chain amino acid sequence

<400> 319

Glu	Val	Gln	Leu	Leu	Glu	Ser	Gly	Gly	Gly	Leu	Val	Gln	Pro	Gly	Gly
1				5					10					15	
Ser	Leu	Arg	Leu	Ser	Cys	Ala	Ala	Ser	Gly	Phe	Thr	Phe	Ser	Leu	Tyr
			20					25					30		
His	Met	Asp	Trp	Val	Arg	Gln	Ala	Pro	Gly	Lys	Gly	Leu	Glu	Trp	Val
		35				40						45			
Ser	Val	Ile	Tyr	Pro	Ser	Gly	Gly	Gly	Thr	Pro	Tyr	Ala	Asp	Ser	Val
	50					55				60					
Lys	Gly	Arg	Phe	Thr	Ile	Ser	Arg	Asp	Asn	Ser	Lys	Asn	Thr	Leu	Tyr
65					70					75				80	
Leu	Gln	Met	Asn	Ser	Leu	Arg	Ala	Glu	Asp	Thr	Ala	Val	Tyr	Tyr	Cys
			85						90					95	
Ala	Arg	Arg	Val	Gly	Tyr	Cys	Ser	Gly	Gly	Ser	Cys	Tyr	Tyr	Tyr	Tyr
			100					105					110		
Tyr	Tyr	Met	Asp	Val	Trp	Gly	Lys	Gly	Thr	Thr	Val	Thr	Val	Ser	Ser
		115				120						125			
Ala	Ser	Thr	Lys	Gly	Pro	Ser	Val	Phe	Pro						
			130				135								

<210> 320
 <211> 14
 <212> PRT
 <213> Unknown

<220>

<223> Light Chain amino acid sequence

<400> 320

Thr	Gly	Thr	Ser	Ser	Asp	Val	Gly	Gly	Tyr	Asn	Tyr	Val	Ser
1				5					10				

<210> 321
 <211> 6
 <212> PRT
 <213> Unknown

<220>

<223> Light Chain amino acid sequence

<400> 321

Asp	Val	Thr	Asn	Arg	Pro
1				5	

<210> 322
 <211> 10
 <212> PRT
 <213> Unknown

<220>

<223> Light Chain amino acid sequence

<400> 322

Val Ser Phe Thr Asn Ser Asn Thr Phe Val
1 5 10

<210> 323

<211> 5

<212> PRT

<213> Unknown

<220>

<223> Heavy Chain amino acid sequence

<400> 323

Leu Tyr His Met Asp
1 5

<210> 324

<211> 17

<212> PRT

<213> Unknown

<220>

<223> Heavy Chain amino acid sequence

<400> 324

Val Ile Tyr Pro Ser Gly Gly Gly Thr Pro Tyr Ala Asp Ser Val Lys
1 5 10 15
Gly

<210> 325

<211> 19

<212> PRT

<213> Unknown

<220>

<223> Heavy Chain amino acid sequence

<400> 325

Arg Val Gly Tyr Cys Ser Gly Gly Ser Cys Tyr Tyr Tyr Tyr Tyr Tyr
1 5 10 15
Met Asp Val

<210> 326

<211> 116

<212> PRT

<213> Unknown

<220>

<223> Light Chain amino acid sequence

<400> 326

Gln Asp Ile Gln Met Thr Gln Ser Pro Ala Thr Leu Ser Val Ser Pro
1 5 10 15

```

Gly Glu Arg Ala Thr Leu Ser Cys Arg Ala Ser Gln Ser Val Arg Ser
      20      25      30
Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro Arg Leu Leu
      35      40      45
Ile Tyr Asp Ala Ser Thr Arg Ala Thr Gly Ile Pro Ala Arg Phe Ser
      50      55      60
Gly Ser Gly Ser Gly Thr Glu Phe Thr Leu Thr Ile Ser Ser Leu Gln
65      70      75      80
Ser Glu Asp Phe Ala Val Tyr Tyr Cys Gln Gln Tyr Asn Asn Trp Pro
      85      90      95
Pro Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys Arg Thr Val Ala
      100      105      110
Ala Pro Ser Val
      115

```

<210> 327

<211> 123

<212> PRT

<213> Unknown

<220>

<223> Heavy Chain amino acid sequence

<400> 327

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Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1      5      10      15
Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Trp Tyr
      20      25      30
Arg Met Asn Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
      35      40      45
Ser Ser Ile Val Pro Ser Gly Gly Tyr Thr Arg Tyr Ala Asp Ser Val
50      55      60
Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65      70      75      80
Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
      85      90      95
Ala Ser Asp Phe Gly Ser Trp Gly Gln Gly Thr Leu Val Thr Val Ser
      100      105      110
Ser Ala Ser Thr Lys Gly Pro Ser Val Phe Pro
      115      120

```

<210> 328

<211> 11

<212> PRT

<213> Unknown

<220>

<223> Light Chain amino acid sequence

<400> 328

```

Arg Ala Ser Gln Ser Val Arg Ser Tyr Leu Ala
1      5      10

```

<210> 329

<211> 7

<212> PRT

<213> Unknown

<220>

<223> Light Chain amino acid sequence

<400> 329

Asp Ala Ser Thr Arg Ala Thr

1

5

<210> 330

<211> 9

<212> PRT

<213> Unknown

<220>

<223> Light Chain amino acid sequence

<400> 330

Gln Gln Tyr Asn Asn Trp Pro Pro Thr

1

5

<210> 331

<211> 5

<212> PRT

<213> Unknown

<220>

<223> Heavy Chain amino acid sequence

<400> 331

Trp Tyr Arg Met Asn

1

5

<210> 332

<211> 17

<212> PRT

<213> Unknown

<220>

<223> Heavy Chain amino acid sequence

<400> 332

Ser Ile Val Pro Ser Gly Gly Tyr Thr Arg Tyr Ala Asp Ser Val Lys

1

5

10

15

Gly

<210> 333

<211> 4

<212> PRT

<213> Unknown

<220>

<223> Heavy Chain amino acid sequence

<400> 333

Asp Phe Gly Ser

1

<210> 334
 <211> 123
 <212> PRT
 <213> Unknown

<220>

<223> Light Chain amino acid sequence

<400> 334

Phe	Tyr	Ser	His	Ser	Ala	Gln	Ser	Glu	Leu	Thr	Gln	Pro	Pro	Ser	Ala
1				5					10					15	
Ser	Gly	Thr	Pro	Gly	Gln	Arg	Val	Thr	Ile	Ser	Cys	Ser	Gly	Ser	Ser
			20					25					30		
Ser	Asn	Ile	Gly	Ser	Asn	Thr	Val	Asn	Trp	Tyr	Gln	Gln	Leu	Pro	Gly
		35					40					45			
Thr	Ala	Pro	Lys	Leu	Leu	Ile	Tyr	Ser	Asn	Asn	Tyr	Arg	Pro	Ser	Gly
	50					55					60				
Val	Pro	Asp	Arg	Phe	Ser	Gly	Ser	Lys	Ser	Gly	Thr	Ser	Ala	Ser	Leu
65					70				75						80
Ala	Ile	Ser	Gly	Leu	Gln	Ser	Asp	Asp	Glu	Ala	Glu	Tyr	Leu	Cys	Ala
				85					90					95	
Ala	Trp	Asp	Asp	Ser	Leu	Asn	Gly	Pro	Val	Phe	Gly	Gly	Gly	Thr	Lys
			100					105					110		
Val	Thr	Val	Leu	Gly	Gln	Pro	Lys	Ala	Ala	Pro					
		115					120								

<210> 335
 <211> 130
 <212> PRT
 <213> Unknown

<220>

<223> Heavy Chain amino acid sequence

<400> 335

Glu	Val	Gln	Leu	Leu	Glu	Ser	Gly	Gly	Gly	Leu	Val	Gln	Pro	Gly	Gly
1				5					10					15	
Ser	Leu	Arg	Leu	Ser	Cys	Ala	Ala	Ser	Gly	Phe	Thr	Phe	Ser	Ser	Tyr
			20					25					30		
Val	Met	Ile	Trp	Val	Arg	Gln	Ala	Pro	Gly	Lys	Gly	Leu	Glu	Trp	Val
		35				40						45			
Ser	Trp	Ile	Ser	Ser	Ser	Gly	Gly	Tyr	Thr	Ser	Tyr	Ala	Asp	Ser	Val
	50					55					60				
Lys	Gly	Arg	Phe	Thr	Ile	Ser	Arg	Asp	Asn	Ser	Lys	Asn	Thr	Leu	Tyr
65					70				75					80	
Leu	Gln	Met	Asn	Ser	Leu	Arg	Ala	Glu	Asp	Thr	Ala	Val	Tyr	Tyr	Cys
			85					90						95	
Ala	Lys	Gly	Pro	Gly	Thr	Arg	Gly	Asp	Tyr	Trp	Gly	Gln	Gly	Thr	Leu
			100					105					110		
Val	Thr	Val	Ser	Ser	Ala	Ser	Thr	Lys	Gly	Pro	Ser	Val	Phe	Pro	Leu
		115					120					125			
Ala	Pro														
	130														

<210> 336
 <211> 13

<212> PRT
 <213> Unknown

<220>
 <223> Light Chain amino acid sequence

<400> 336
 Ser Gly Ser Ser Ser Asn Ile Gly Ser Asn Thr Val Asn
 1 5 10

<210> 337
 <211> 5
 <212> PRT
 <213> Unknown

<220>
 <223> Heavy Chain amino acid sequence

<400> 337
 Ser Tyr Val Met Ile
 1 5

<210> 338
 <211> 17
 <212> PRT
 <213> Unknown

<220>
 <223> Heavy Chain amino acid sequence

<400> 338
 Trp Ile Ser Ser Ser Gly Gly Tyr Thr Ser Tyr Ala Asp Ser Val Lys
 1 5 10 15
 Gly

<210> 339
 <211> 8
 <212> PRT
 <213> Unknown

<220>
 <223> Heavy Chain amino acid sequence

<400> 339
 Gly Pro Gly Thr Arg Gly Asp Tyr
 1 5

<210> 340
 <211> 123
 <212> PRT
 <213> Unknown

<220>
 <223> Light Chain amino acid sequence

<400> 340


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Phe Tyr Ser His Ser Ala Gln Ser Val Leu Thr Gln Pro Pro Ser Ala
 1           5           10           15
Ser Ala Thr Pro Gly Gln Arg Val Thr Phe Ser Cys Ser Gly Ser Ser
          20           25           30
Ser Asn Ile Gly Ser Asn Ala Val Asn Trp Tyr His Gln Leu Pro Gly
          35           40           45
Thr Ala Pro Lys Leu Leu Ile Tyr His Asn Asn Gln Arg Pro Ser Gly
          50           55           60
Val Pro Asp Arg Phe Ser Gly Ser Lys Ser Gly Thr Ser Ala Ser Leu
65           70           75           80
Ala Ile Ser Gly Leu Gln Ser Glu Asp Glu Ala Asp Tyr Tyr Cys Ala
          85           90           95
Ala Trp Asp Asp Ser Leu His Gly Tyr Val Phe Gly Pro Gly Thr Lys
          100          105          110
Val Thr Val Leu Gly Gln Pro Lys Ala Asn Pro
          115          120

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<210> 341
 <211> 131
 <212> PRT
 <213> Unknown

<220>
 <223> Heavy Chain amino acid sequence

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<400> 341
Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
 1           5           10           15
Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ile Tyr
          20           25           30
Pro Met Asn Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
          35           40           45
Ser Gly Ile Ser Pro Ser Gly Gly Tyr Thr Gly Tyr Ala Asp Ser Val
          50           55           60
Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65           70           75           80
Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
          85           90           95
Ala Arg Gly Gly Ile Ser Trp Phe Met Asp Tyr Trp Gly Gln Gly Thr
          100          105          110
Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val Phe Pro
          115          120          125
Leu Ala Pro
          130

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<210> 342
 <211> 13
 <212> PRT
 <213> Unknown

<220>
 <223> Light Chain amino acid sequence

```

<400> 342
Ser Gly Ser Ser Ser Asn Ile Gly Ser Asn Ala Val Asn
 1           5           10

```

<210> 343
 <211> 7
 <212> PRT
 <213> Unknown

<220>
 <223> Light Chain amino acid sequence

<400> 343
 His Asn Asn Gln Arg Pro Ser
 1 5

<210> 344
 <211> 11
 <212> PRT
 <213> Unknown

<220>
 <223> Light Chain amino acid sequence

<400> 344
 Ala Ala Trp Asp Asp Ser Leu His Gly Tyr Val
 1 5 10

<210> 345
 <211> 5
 <212> PRT
 <213> Unknown

<220>
 <223> Heavy Chain amino acid sequence

<400> 345
 Ile Tyr Pro Met Asn
 1 5

<210> 346
 <211> 17
 <212> PRT
 <213> Unknown

<220>
 <223> Heavy Chain amino acid sequence

<400> 346
 Gly Ile Ser Pro Ser Gly Gly Tyr Thr Gly Tyr Ala Asp Ser Val Lys
 1 5 10 15
 Gly

<210> 347
 <211> 9
 <212> PRT
 <213> Unknown

<220>
 <223> Heavy Chain amino acid sequence

<400> 347

Gly Gly Ile Ser Trp Phe Met Asp Tyr

1

5

<210> 348

<211> 369

<212> DNA

<213> Unknown

<220>

<223> Light Chain nucleic acid sequence

<400> 348

ttctattctc	acagtgcaca	gagcgtcttg	actcagccac	cctcagcgtc	tgcgaccccc	60
gggcagaggg	tcaccttctc	ttgttctgga	agcagctcca	acatcggaag	taatgctgta	120
aactggtacc	atcagctccc	aggaacggcc	cccaaactcc	tcattctatca	taataatcag	180
cgaccctcag	gggtccctga	ccgattctct	ggctccaagt	ctggcacctc	agcctccctg	240
gccatcagtg	ggctccagtc	tgaggatgag	gctgattatt	actgtgcagc	atgggatgac	300
agcctgcatg	gttatgtctt	cggacctggg	accaaggtca	ccgtcctagg	tcagcccaag	360
gccaacccc						369

<210> 349

<211> 393

<212> DNA

<213> Unknown

<220>

<223> Heavy Chain nucleic acid sequence

<400> 349

gaagttcaat	tgtagagtc	tggtggcgg	cttggtcagc	ctgggtggtc	tttacgtctt	60
tcttgcgctg	cttccggatt	cactttctct	atttacccta	tgaattgggt	tcgccaagct	120
cctggtaaa	gtttgagtg	ggtttctgg	atctctcctt	ctgggtggta	tactggttat	180
gctgactccg	ttaaaggctg	cttcaactat	tctagagaca	actctaagaa	tactctctac	240
ttgcagatga	acagcttaag	ggctgaggac	actgcagctc	actattgtgc	gagagggggc	300
atcagctgg	ttatggacta	ctggggccag	ggaaccctgg	tcaccgtctc	aagcgccctc	360
accaagggcc	catcggtctt	cccgttagca	ccc			393

<210> 350

<211> 378

<212> DNA

<213> Unknown

<220>

<223> Light Chain nucleic acid sequence

<400> 350

ttctattctc	acagtgcaca	gagcgtcttg	actcagcctc	gctcagtgtc	cgggtctcct	60
ggacagtcag	tcaccatctc	ctgcactgga	accagtagtg	atgttggtgc	tagttataag	120
tttgtctcct	ggtaccaact	aaagccaggc	aaagcccca	aactcatgct	ttttaatgtc	180
cgtgagcggc	cctcaggggt	ccctgatcgc	ttttctgggt	ccaagtccgg	caacacggcc	240
tccctgacca	tctctgggct	ccaggctgag	gatgaggctg	actattactg	ctgttcctat	300
gcacgcggcc	agactttctc	ttatgtcttc	ggaggtggga	ccacggtcac	cgtcctagg	360
cagcccaagg	ccaacccc					378

<210> 351

<211> 402
 <212> DNA
 <213> Unknown

<220>
 <223> Heavy Chain nucleic acid sequence

```
<400> 351
gaagttcaat tgtagagtc tggtagcggg cttgttcagc ctggtgggtc tttacgtctt      60
tcttgcgctg cttccggatt cactttctct cgttactcta tgggggtgggt tcgccaagct      120
cctggtaaag gtttggagtg ggtttcttct atccgtcctt ctggtggcta tactcgttat      180
gctgactccg ttaaaggctg cttcactatc tctagagaca actctaagaa tactctctac      240
ttgcagatga acagcttaag ggctgaggac actgcagtct actattgtgc gaaagatctg      300
gagtatagca gtggctggtc atttgactac tggggccagg gaaccctggg caccgtctca      360
agcgctcca ccaagggcc atcggtcttc ccgctagcac cc                                402
```

<210> 352
 <211> 126
 <212> PRT
 <213> Unknown

<220>
 <223> Light Chain amino acid sequence

```
<400> 352
Phe Tyr Ser His Ser Ala Gln Ser Val Leu Thr Gln Pro Arg Ser Val
 1          5          10          15
Ser Gly Ser Pro Gly Gln Ser Val Thr Ile Ser Cys Thr Gly Thr Ser
 20          25          30
Ser Asp Val Gly Ala Ser Tyr Lys Phe Val Ser Trp Tyr Gln Leu Lys
 35          40          45
Pro Gly Lys Ala Pro Lys Leu Met Leu Phe Asn Val Arg Glu Arg Pro
 50          55          60
Ser Gly Val Pro Asp Arg Phe Ser Gly Ser Lys Ser Gly Asn Thr Ala
 65          70          75          80
Ser Leu Thr Ile Ser Gly Leu Gln Ala Glu Asp Glu Ala Asp Tyr Tyr
 85          90          95
Cys Cys Ser Tyr Ala Arg Gly Gln Thr Phe Ser Tyr Val Phe Gly Gly
100          105          110
Gly Thr Thr Val Thr Val Leu Gly Gln Pro Lys Ala Asn Pro
115          120          125
```

<210> 353
 <211> 134
 <212> PRT
 <213> Unknown

<220>
 <223> Heavy Chain amino acid sequence

```
<400> 353
Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
 1          5          10          15
Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Arg Tyr
 20          25          30
Ser Met Gly Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
 35          40          45
```

Ser Ser Ile Arg Pro Ser Gly Gly Tyr Thr Arg Tyr Ala Asp Ser Val
 50 55 60
 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
 65 70 75 80
 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
 85 90 95
 Ala Lys Asp Leu Glu Tyr Ser Ser Gly Trp Ser Phe Asp Tyr Trp Gly
 100 105 110
 Gln Gly Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser
 115 120 125
 Val Phe Pro Leu Ala Pro
 130

<210> 354
 <211> 12
 <212> PRT
 <213> Unknown

<220>
 <223> Light Chain amino acid sequence

<400> 354
 Cys Ser Tyr Ala Arg Gly Gln Thr Phe Ser Tyr Val
 1 5 10

<210> 355
 <211> 5
 <212> PRT
 <213> Unknown

<220>
 <223> Heavy Chain amino acid sequence

<400> 355
 Arg Tyr Ser Met Gly
 1 5

<210> 356
 <211> 17
 <212> PRT
 <213> Unknown

<220>
 <223> Heavy Chain amino acid sequence

<400> 356
 Ser Ile Arg Pro Ser Gly Gly Tyr Thr Arg Tyr Ala Asp Ser Val Lys
 1 5 10 15
 Gly

<210> 357
 <211> 12
 <212> PRT
 <213> Unknown

<220>

<223> Heavy Chain amino acid sequence

<400> 357

Asp	Leu	Glu	Tyr	Ser	Ser	Gly	Trp	Ser	Phe	Asp	Tyr
1					5					10	

<210> 358

<211> 11

<212> PRT

<213> Artificial Sequence

<220>

<223> Exemplary motif

<221> VARIANT

<222> 4

<223> Xaa = Gln or Arg

<221> VARIANT

<222> 5

<223> Xaa = Asp, Gly, Arg, or Ser

<221> VARIANT

<222> 6

<223> Xaa = Val or Ile

<221> VARIANT

<222> 7

<223> Xaa = Arg, Ser or Asn

<221> VARIANT

<222> 8

<223> Xaa = Asn, Arg, His, Ser or Thr

<221> VARIANT

<222> (9)...(0)

<223> Xaa = Tyr, Asp, Glu, Trp, Asn or Ser

<221> VARIANT

<222> (10)...(0)

<223> Xaa = Leu, Val, or Tyr

<221> VARIANT

<222> (11)...(0)

<223> Xaa = Ala, Gly, Asn or Leu

<400> 358

Arg	Ala	Ser	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa
1				5						10	

<210> 359

<211> 10

<212> PRT

<213> Artificial Sequence

<220>

<223> Exemplary motif

<221> VARIANT
 <222> 5, 7, 8
 <223> Xaa = any amino acid, e.g., a hydrophilic amino acid

<221> VARIANT
 <222> 6
 <223> Xaa = Val or Ile

<221> VARIANT
 <222> 9
 <223> Xaa = Tyr, Asp, Glu, Trp, Asn or Ser

<221> VARIANT
 <222> 10
 <223> Xaa = is hydrophobic, or aliphatic

<400> 359
 Arg Ala Ser Gln Xaa Xaa Xaa Xaa Xaa Xaa
 1 5 10

<210> 360
 <211> 13
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Exemplary motif

<221> VARIANT
 <222> 8
 <223> Xaa = Gly, Glu, Asp or Ala

<221> VARIANT
 <222> 9
 <223> Xaa = Ser, Arg or Val

<221> VARIANT
 <222> 10
 <223> Xaa = Asn or Tyr

<221> VARIANT
 <222> 11
 <223> Xaa = Thr, Leu, Phe or Asp

<221> VARIANT
 <222> 13
 <223> Xaa = Tyr or Thr

<400> 360
 Ser Gly Ser Ser Ser Asn Ile Xaa Xaa Xaa Xaa Val Xaa
 1 5 10

<210> 361
 <211> 14
 <212> PRT

<213> Artificial Sequence

<220>

<223> Exemplary motif

<221> VARIANT

<222> 7

<223> Xaa = Ile or Val

<221> VARIANT

<222> 9

<223> Xaa = Asp, Gly or Tyr

<221> VARIANT

<222> 11

<223> Xaa = Asn, Glu or Asp

<400> 361

Thr Gly Thr Ser Ser Asp Xaa Gly Xaa Tyr Xaa Tyr Val Ser
1 5 10

<210> 362

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> Exemplary motif

<221> VARIANT

<222> 1

<223> Xaa = Ser or Thr

<221> VARIANT

<222> 2, 3

<223> Xaa = Asp or Asn

<400> 362

Xaa Xaa Xaa Gln Arg Pro Ser
1 5

<210> 363

<211> 6

<212> PRT

<213> Artificial Sequence

<220>

<223> Exemplary motif

<221> VARIANT

<222> 4

<223> Xaa = Ser or Thr

<221> VARIANT

<222> 5

<223> Xaa = Leu or Arg

<221> VARIANT

<222> 6

<223> Xaa = Gln or Ala

<400> 363

Gly Ala Ser Xaa Xaa Xaa

1

5

<210> 364

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Exemplary motif

<221> VARIANT

<222> 1

<223> Xaa = Gln or Leu

<221> VARIANT

<222> 3

<223> Xaa = any amino acid or is hydrophilic, Ala, or Gly,

<221> VARIANT

<222> 4, 5

<223> Xaa = any amino acid or is hydrophilic

<221> VARIANT

<222> 6

<223> Xaa = aromatic, Thr , Arg or Lys

<221> VARIANT

<222> 8

<223> Xaa is hydrophobic

<400> 364

Xaa Gln Xaa Xaa Xaa Xaa Pro Xaa

1

5

<210> 365

<211> 9

<212> PRT

<213> Artificial Sequence

<220>

<223> Exemplary motif

<221> VARIANT

<222> 4, 5

<223> Xaa = any amino acid

<221> VARIANT

<222> 6

<223> Xaa = hydrophobic (e.g., aromatic)

<221> VARIANT

<222> 8

<223> Xaa = Pro, Leu or Arg

<400> 365

Gln Gln Tyr Xaa Xaa Xaa Pro Xaa Thr
1 5

<210> 366

<211> 10

<212> PRT

<213> Artificial Sequence

<220>

<223> Exemplary motif

<221> VARIANT

<222> 9

<223> Xaa = hydrophobic

<400> 366

Ala Trp Asp Asp Ser Leu Ser Gly Xaa Val
1 5 10

<210> 367

<211> 10

<212> PRT

<213> Artificial Sequence

<220>

<223> Exemplary motif

<221> VARIANT

<222> 9

<223> Xaa = Val or Trp

<400> 367

Ala Trp Asp Asp Ser Leu Ser Gly Xaa Val
1 5 10

<210> 368

<211> 11

<212> PRT

<213> Artificial Sequence

<220>

<223> Exemplary motif

<221> VARIANT

<222> 2

<223> Xaa = Ala or Thr

<221> VARIANT

<222> 5

<223> Xaa = Asp, Asn, Glu or Gln

<221> VARIANT

<222> 6
 <223> Xaa = Ser or Thr

<221> VARIANT
 <222> 8
 <223> Xaa = Ser, Arg or Thr

<221> VARIANT
 <222> 10
 <223> Xaa = Val or trp

<400> 368
 Ala Xaa Trp Asp Xaa Xaa Leu Xaa Gly Xaa Val
 1 5 10

<210> 369
 <211> 4
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Exemplary motif

<221> VARIANT
 <222> 2
 <223> Xaa = any amino acid , Trp, Asp, Lys, Thr, Arg,
 His or Pro

<221> VARIANT
 <222> 4
 <223> Xaa = Asn, Trp, Asp, Glu, Pro, Thr, Arg, Ser, Val
 or Phe

<400> 369
 Tyr Xaa Met Xaa
 1

<210> 370
 <211> 5
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Exemplary motif

<221> VARIANT
 <222> 1
 <223> Xaa = aromatic

<221> VARIANT
 <222> 3
 <223> Xaa = any amino acid

<221> VARIANT
 <222> 5
 <223> Xaa = Asn, Trp, Asp, Glu, Pro, Thr, Ser, Val or
 Phe

<400> 370

Xaa Tyr Xaa Met Xaa

1 5

<210> 371

<211> 4

<212> PRT

<213> Artificial Sequence

<220>

<223> Exemplary motif

<221> VARIANT

<222> 3

<223> Xaa = any amino acid, Trp, His or Thr

<400> 371

Trp Tyr Xaa Met

1

<210> 372

<211> 4

<212> PRT

<213> Artificial Sequence

<220>

<223> Exemplary motif

<221> VARIANT

<222> 3

<223> Xaa = any amino acid

<400> 372

Gln Tyr Xaa Met

1

<210> 373

<211> 6

<212> PRT

<213> Artificial Sequence

<220>

<223> Exemplary motif

<221> VARIANT

<222> 2

<223> Xaa = any amino acid, hydrophobic or Val, Tyr,
Trp, Arg, Ser, or Gly

<221> VARIANT

<222> 3

<223> Xaa = Pro or Ser

<400> 373

Ile Xaa Xaa Ser Gly Gly

1 5

<210> 374
 <211> 8
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Exemplary motif

<221> VARIANT
 <222> 2, 7
 <223> Xaa = any amino acid

<221> VARIANT
 <222> 3
 <223> Xaa = Pro or Ser

<400> 374
 Ile Xaa Xaa Ser Gly Gly Xaa Thr
 1 5

<210> 375
 <211> 8
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Exemplary motif

<221> VARIANT
 <222> 2
 <223> Xaa = Ser, Val, Tyr, Trp, Arg or Gly

<221> VARIANT
 <222> 3
 <223> xaa = Pro or Ser

<221> VARIANT
 <222> 7
 <223> Xaa = Gly, Lys, Leu, Arg, His, Phe, Tyr, Thr, Gly,
 Gln, Asp, Met, Ile or Asn

<400> 375
 Ile Xaa Xaa Ser Gly Gly Xaa Thr
 1 5

<210> 376
 <211> 16
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Exemplary motif

<221> VARIANT
 <222> 2
 <223> Xaa = Ser, Val, Tyr, Trp, Arg or Gly

<221> VARIANT

<222> 3

<223> Xaa = Pro or Ser

<221> VARIANT

<222> 7, 9

<223> Xaa = any amino acid

<400> 376

Ile	Xaa	Xaa	Ser	Gly	Gly	Xaa	Thr	Xaa	Tyr	Ala	Asp	Ser	Val	Lys	Gly
1				5					10					15	

<210> 377

<211> 4

<212> PRT

<213> Artificial Sequence

<220>

<223> Exemplary motif

<221> VARIANT

<222> 1, 2

<223> Xaa = Ser or Gly

<400> 377

Xaa	Xaa	Trp	Tyr
1			

<210> 378

<211> 5

<212> PRT

<213> Artificial Sequence

<220>

<223> Exemplary motif

<221> VARIANT

<222> 3

<223> Xaa = Ser or Gly

<400> 378

Ser	Ser	Xaa	Trp	Tyr
1			5	

<210> 379

<211> 6

<212> PRT

<213> Artificial Sequence

<220>

<223> Exemplary motif

<221> VARIANT

<222> 1

<223> Xaa = Arg, His, Trp or Tyr

<400> 379
 Xaa Tyr Tyr Tyr Gly Met
 1 5

<210> 380
 <211> 8
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Exemplary motif

<221> VARIANT
 <222> 1
 <223> Xaa = Tyr, Ser or Gly

<221> VARIANT
 <222> 2
 <223> Xaa = Arg, His, Trp or Tyr

<400> 380
 Xaa Xaa Tyr Tyr Tyr Gly Met Asp
 1 5

<210> 381
 <211> 11
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Exemplary motif

<221> VARIANT
 <222> 1
 <223> Xaa = Ala, Gly, Gln, Ser or Val

<221> VARIANT
 <222> 2
 <223> Xaa = Ala, Thr or Ser

<221> VARIANT
 <222> 3
 <223> Xaa = aromatic

<221> VARIANT
 <222> 4
 <223> Xaa = any amino acid, or Glu, Asp, Arg, Thr or Ser

<221> VARIANT
 <222> 5
 <223> Xaa = any amino acid, or Asp, Asn, Gln, Lys, Arg
 or Ser

<221> VARIANT
 <222> (7)...(0)
 <223> Xaa = any amino acid, or Ser, Leu, Thr or Asn

<221> VARIANT
 <222> (6)...(0)
 <223> Xaa = Ser, Thr, Gly or Ala

<221> VARIANT
 <222> (8)...(0)
 <223> Xaa = Ser, Thr, Arg or Gly

<221> VARIANT
 <222> (9)...(0)
 <223> Xaa = Gly, Pro, Asn or Phe

<221> VARIANT
 <222> (10)...(0)
 <223> Xaa = any amino acid

<400> 381
 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Val
 1 5 10

<210> 382
 <211> 7
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Exemplary motif

<221> VARIANT
 <222> 1
 <223> Xaa = Ser or Thr

<221> VARIANT
 <222> 2, 3, 4
 <223> Xaa = hydrophilic

<221> VARIANT
 <222> 5
 <223> Xaa = Leu, Arg or Asn

<221> VARIANT
 <222> 6
 <223> Xaa = pro, Arg or Gln

<400> 382
 Xaa Xaa Xaa Xaa Xaa Xaa Ser
 1 5

<210> 383
 <211> 5
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Exemplary motif

<221> VARIANT

<222> 3
 <223> Xaa = Ser or Gly

<221> VARIANT
 <222> 5
 <223> Xaa = Ser or Tyr

<400> 383
 Ser Ser Xaa Trp Xaa
 1 5

<210> 384
 <211> 7
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Exemplary motif

<221> VARIANT
 <222> 1
 <223> Xaa = Ala, Asp, Glu or Gly

<221> VARIANT
 <222> 2
 <223> Xaa = Ala, Val, Asp, Asn or Glu

<221> VARIANT
 <222> 3
 <223> Xaa = Ala, Ser, Thr, Asn or Val

<221> VARIANT
 <222> 4
 <223> Xaa = Ser, Thr, Asn or Gln

<221> VARIANT
 <222> 5
 <223> Xaa = Leu, Arg or Asn

<221> VARIANT
 <222> (6)...(0)
 <223> Xaa = Ala, Gln, Pro or Arg

<221> VARIANT
 <222> (7)...(0)
 <223> Xaa = Thr, Phe, Ser, Lys or Pro

<400> 384
 Xaa Xaa Xaa Xaa Xaa Xaa
 1 5

<210> 385
 <211> 7
 <212> PRT
 <213> Artificial Sequence

<220>

<223> Exemplary motif

<221> VARIANT

<222> 1

<223> Xaa = Ala, Asp, Glu, Asn or Gly

<221> VARIANT

<222> 2

<223> Xaa = Ala, Val, Asp, Asn or Glu

<221> VARIANT

<222> 3

<223> Xaa = Ala, Ser, Thr, Arg, Asn or Val

<221> VARIANT

<222> 4

<223> Xaa = Ser, Thr, Asn or Gln

<221> VARIANT

<222> 5

<223> Xaa = Leu, Arg or Asn

<221> VARIANT

<222> (6)...(0)

<223> Xaa = Ala, Gln, Pro or Arg

<221> VARIANT

<222> (70)...(0)

<223> Xaa = Thr, Phe, Ser, Lys or Pro

<400> 385

Xaa Xaa Xaa Xaa Xaa Xaa Xaa

1

5

<210> 386

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> Exemplary motif

<221> VARIANT

<222> 1

<223> Xaa = Ala, Asp or Glu

<221> VARIANT

<222> 2

<223> Xaa = Ala or Val

<221> VARIANT

<222> 3

<223> Xaa = Ala, Ser or Thr

<221> VARIANT

<222> 4

<223> Xaa = Ser or Thr

<221> VARIANT

<222> 5

<223> Xaa = Leu or Arg

<221> VARIANT

<222> (6)...(0)

<223> Xaa = Ala or Gln

<221> VARIANT

<222> (7)...(0)

<223> Xaa = Thr, Phe, Ser or Lys

<400> 386

Xaa Xaa Xaa Xaa Xaa Xaa Xaa

1

5

<210> 387

<211> 14

<212> PRT

<213> Artificial Sequence

<220>

<223> Exemplary motif

<221> VARIANT

<222> 7

<223> Xaa = Ile or Val

<221> VARIANT

<222> 9

<223> Xaa = Ala, Asp, Gly or Tyr

<221> VARIANT

<222> 11

<223> Xaa = Asn, Lys, Glu or Asp

<221> VARIANT

<222> 12

<223> Xaa = Tyr or Phe

<400> 387

Thr Gly Thr Ser Ser Asp Xaa Gly Xaa Tyr Xaa Xaa Val Ser

1

5

10

<210> 388

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> Exemplary motif

<221> VARIANT

<222> 1

<223> Xaa = Asn, Ser or Thr

<221> VARIANT
 <222> 2, 3, 4
 <223> Xaa = hydrophilic

<221> VARIANT
 <222> 5
 <223> Xaa = Leu, Arg or Asn

<221> VARIANT
 <222> 6
 <223> Xaa = Pro, Arg or Gln

<400> 388
 Xaa Xaa Xaa Xaa Xaa Xaa Ser
 1 5

<210> 389
 <211> 13
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Exemplary motif

<221> VARIANT
 <222> 8
 <223> Xaa = Gly, Glu, Asp or Ala

<221> VARIANT
 <222> 9
 <223> Xaa = Ser, Arg or Val

<221> VARIANT
 <222> 10
 <223> Xaa = Ala, Asn or Tyr

<221> VARIANT
 <222> 11
 <223> Xaa = Thr, leu, Phe or Asp

<221> VARIANT
 <222> 13
 <223> Xaa = Asn, Tyr or Thr

<400> 389
 Ser Gly Ser Ser Ser Asn Ile Xaa Xaa Xaa Xaa Val Xaa
 1 5 10

<210> 390
 <211> 6
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Exemplary motif

<221> VARIANT

<222> 1
 <223> Xaa = his, Ser or Thr

<221> VARIANT
 <222> 2, 3
 <223> Xaa = Asp or Asn

<221> VARIANT
 <222> 4
 <223> Xaa = Gln or Tyr

<400> 390
 Xaa Xaa Xaa Xaa Arg Pro
 1 5

<210> 391
 <211> 11
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Exemplary motif

<221> VARIANT
 <222> 2
 <223> Xaa = Ala or Thr

<221> VARIANT
 <222> 5
 <223> Xaa = Asp, Asn, Glu or Gln

<221> VARIANT
 <222> 6
 <223> Xaa = Ser or Thr

<221> VARIANT
 <222> 8
 <223> Xaa = any anini acid, e.g. Ser, Arg, Thr, His or
 Asn

<221> VARIANT
 <222> 10
 <223> Xaa = any amino acid, e.g., hydrophobic, e.g.,
 Val, Tyr or Trp

<400> 391
 Ala Xaa Trp Asp Xaa Xaa Leu Xaa Gly Xaa Val
 1 5 10

<210> 392
 <211> 8
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Exemplary motif

<221> VARIANT
 <222> 1
 <223> Xaa = Asn, Gln, Arg or Lys

<221> VARIANT
 <222> 2
 <223> Xaa = hydrophilic, Ala or Gly

<221> VARIANT
 <222> 3
 <223> Xaa = Aliphatic

<221> VARIANT
 <222> 4, 5
 <223> Xaa = hydrophilic

<221> VARIANT
 <222> 6
 <223> Xaa = any amino acid, or aromatic or hydrophilic

<221> VARIANT
 <222> (7)...(0)
 <223> Xaa = hydrophobic

<400> 392
 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
 1 5

<210> 393
 <211> 10
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Exemplary motif

<221> VARIANT
 <222> 1
 <223> Xaa = Thr or Ser

<221> VARIANT
 <222> 2
 <223> Xaa = Asp or Glu

<221> VARIANT
 <222> 3
 <223> Xaa = Aliphatic

<221> VARIANT
 <222> 5
 <223> Xaa = hydrophilic or Gly

<221> VARIANT
 <222> 7
 <223> Xaa = hydrophilic, or Asn, Glu, Asp or Gln

<400> 393

Xaa Xaa Xaa Gly Xaa Tyr Xaa Xaa Xaa Xaa
1 5 10

<210> 394

<211> 4

<212> PRT

<213> Artificial Sequence

<220>

<223> Exemplary motif

<400> 394

Asp Phe Gly Ser

1